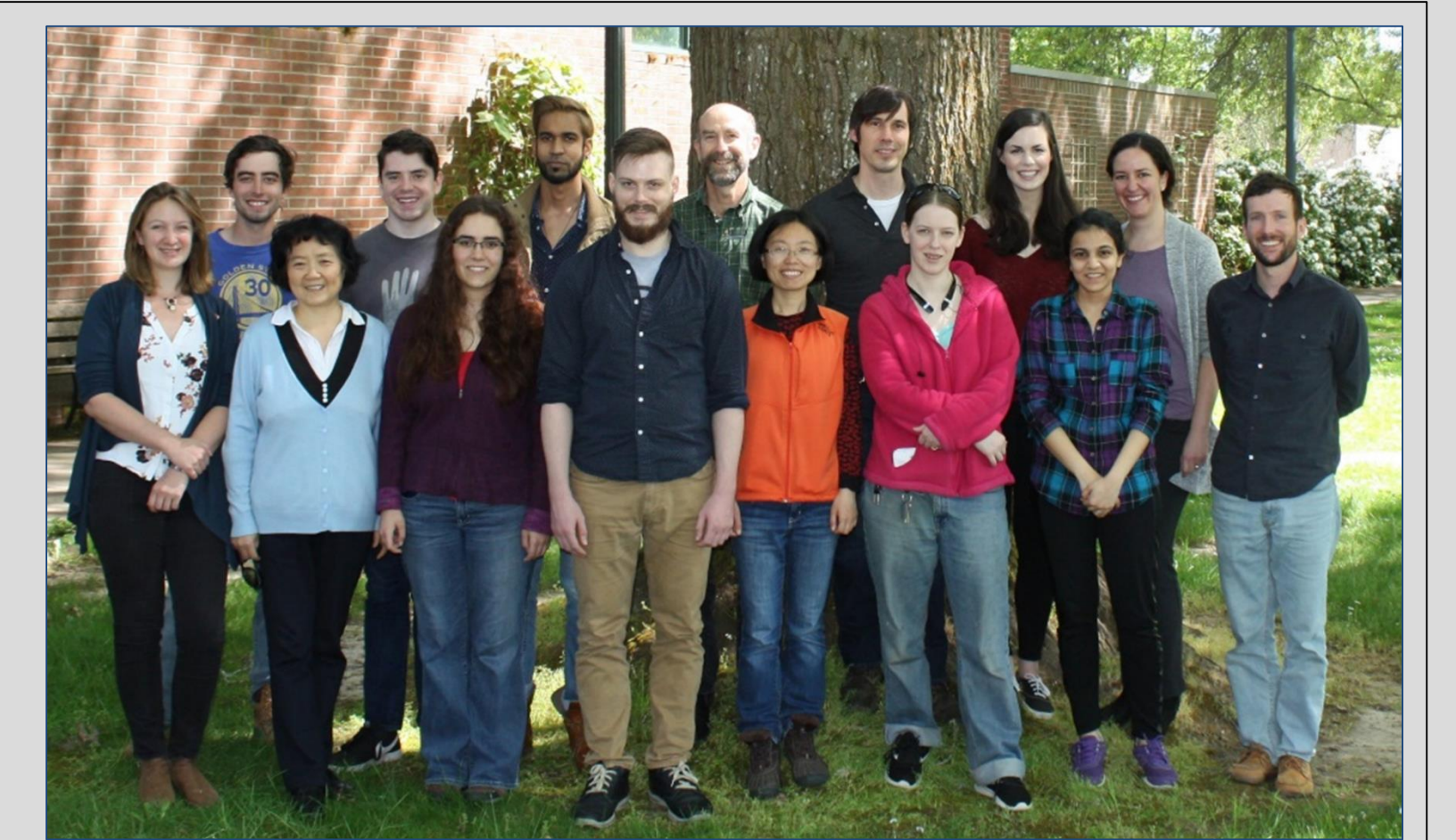


Toward Optimization of *in vitro* Regeneration and Transformation in Wild Black Cottonwood (*Populus trichocarpa*)

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Summary

As part of a Genome Wide Association Study (GWAS) to identify genes that modify the amenability of poplars (*Populus*) to *in vivo* and *in vitro* regeneration and transformation, we studied a wide variety of media components, hormones, and transformation procedures in a sample of resequenced wild cottonwood genotypes.

We analyzed the effects of 36 permutations of MS basal media [factorial combinations of NH₄NO₃, KNO₃, and MESO (CaCl₂, MgSO₄ and KH₂PO₄) levels], resulting in the identification of five media that gave consistent and high rates of organogenesis. When variable levels of sucrose (1.5, 2.0, 2.5, and 3.0%) in the selected MS media were studied, the highest sucrose level supported the strongest callus and shoot regeneration from all genotypes.

We investigated several factors that affect *Agrobacterium*-mediated transformation in three genotypes using the 2X-35S promoter driving an enhanced GFP gene (in Pc2300 backbone). Acetosyringone (AS) increased transient and stable transformation. The rate of recovery of transgenic plants was higher when geneticin was the selection agent instead of kanamycin.

The relationship of transformation rate to duration (0 to 21 days) of time on callus induction medium (CIM) with geneticin (after co-cultivation) was examined, and the optimal durations varied widely among genotypes. The best transformation rate from the genome sequenced clone *P. trichocarpa* Nisqally-1 was 6.3%, and occurred when there was only two days of co-cultivation on CIM (with AS) before explants were moved to shoot induction medium (SIM). Direct regeneration (no CIM phase) gave a high rate of shoot regeneration from many clones but almost all regenerants were escapes. Customizing the CIM:SIM period for individual genotypes seems to be an important step in transformation.

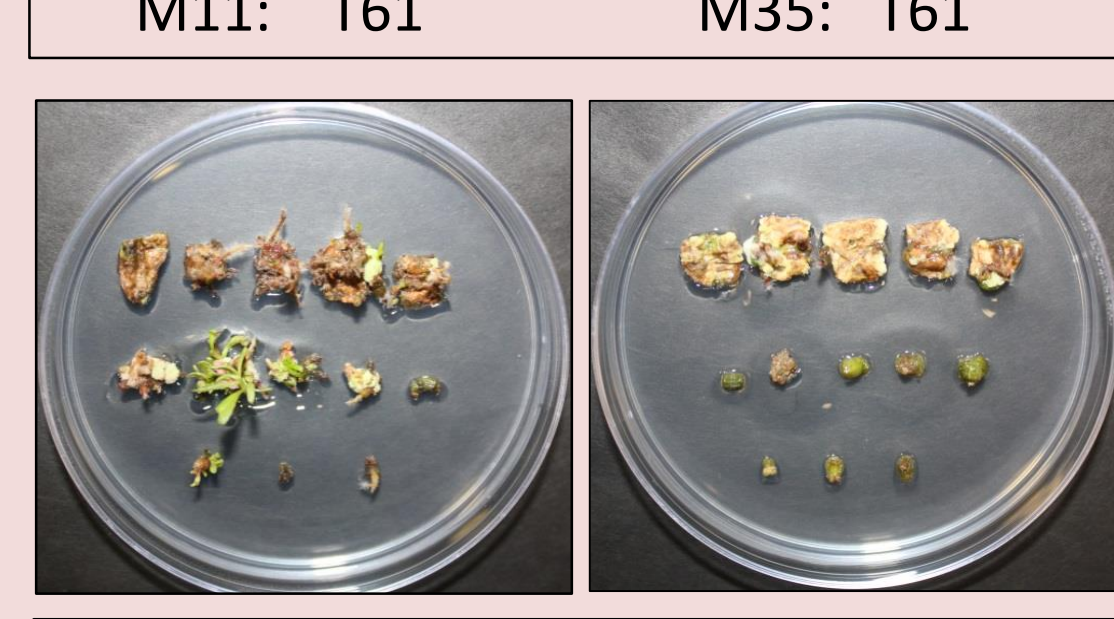
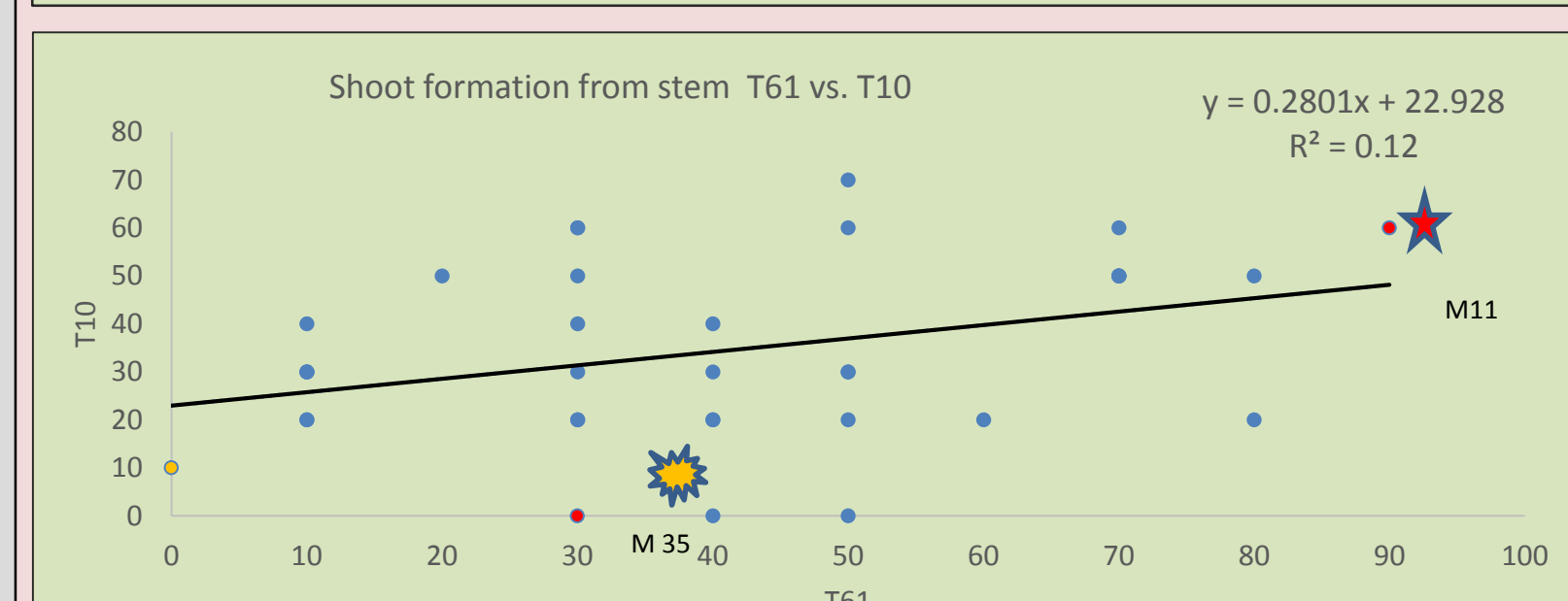
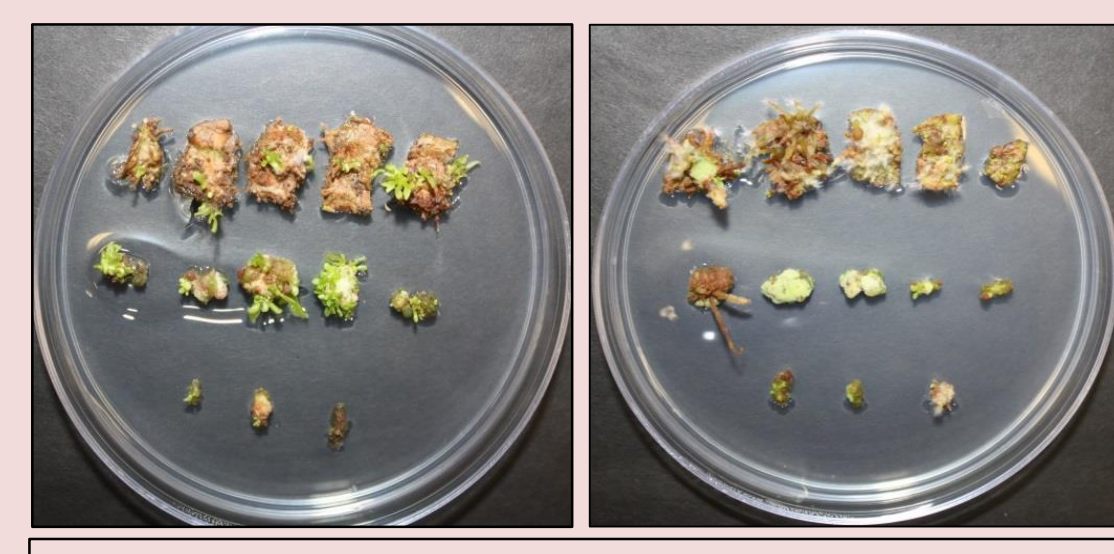
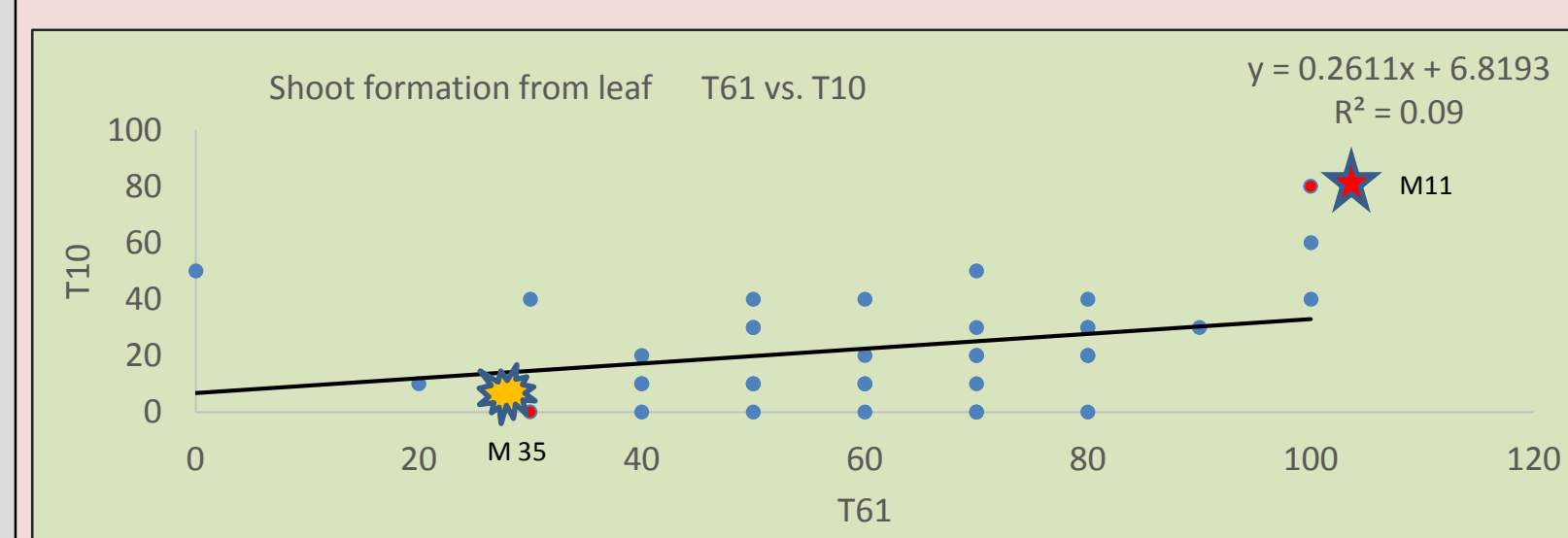
Basal medium screening for shoot regeneration

- 36 basal media were screened (0.25X, 0.5X, 1X, 1.5X of MS in NH₄NO₃, KNO₃, Mesos)
- Leaf, stem and petiole explants were tested
- 12 explants per plate, 2 plates per genotype, studied
- Explants were cultured on callus induction medium (CIM) for 20d in dark and the basal medium was supplemented with 2μM 2iP and 10μM NAA
- Explants were transferred onto shoot induction medium (SIM) for 42 days under light and supplemented with 0.6μM TDZ

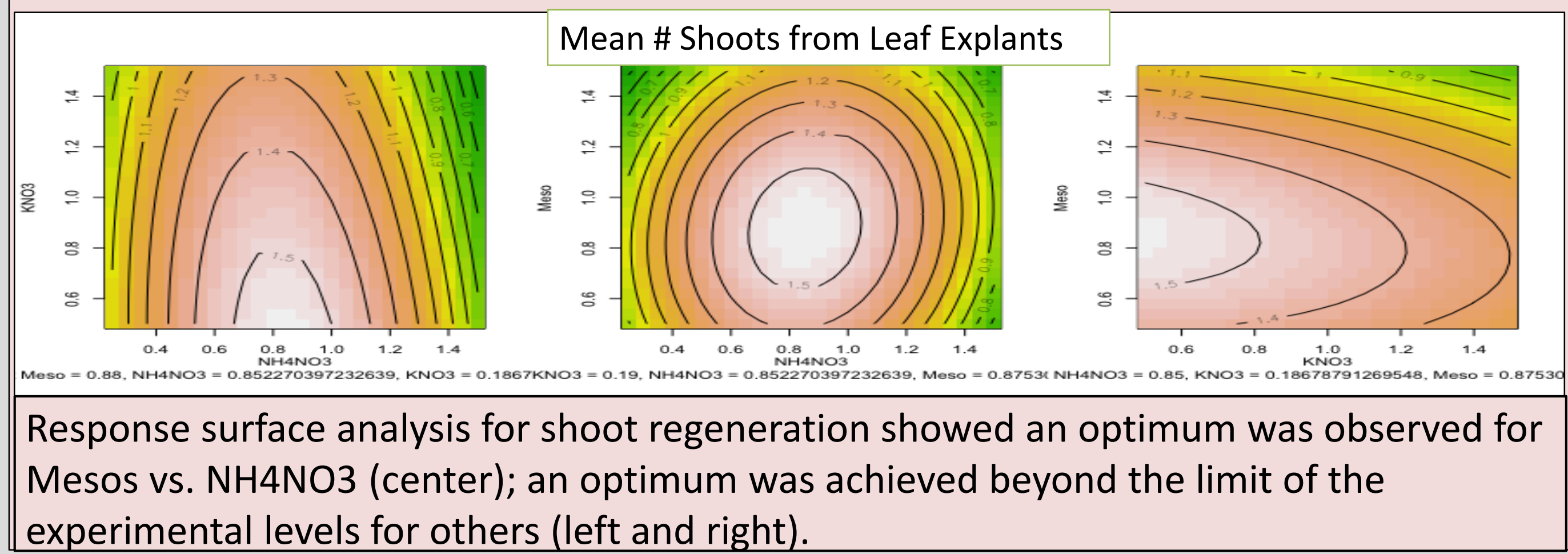
Table 1. List of Optimization Table (X is the amount relative to MS)

Media	NH4NO3	KNO3	Meso	CaCl2	MgSO4	KH2PO4	Micro-nutrients	Fe-EDTA
M11	0.5X	1X	0.5X	1X	1X			
M19	1X	0.5X	0.5X	1X	1X			
M23	1X	1X	1X	1X	1X			
M24	1X	1.5X	1X	1X	1X			
M28	1.5X	0.5X	0.5X	1X	1X			

We identified media that supported high shoot regeneration from diverse clones and explant types



M11 (low salt) gave the best regeneration from both leaf and stem in the two clones, whereas M35 and M30 (high salt) gave poor responses in both.

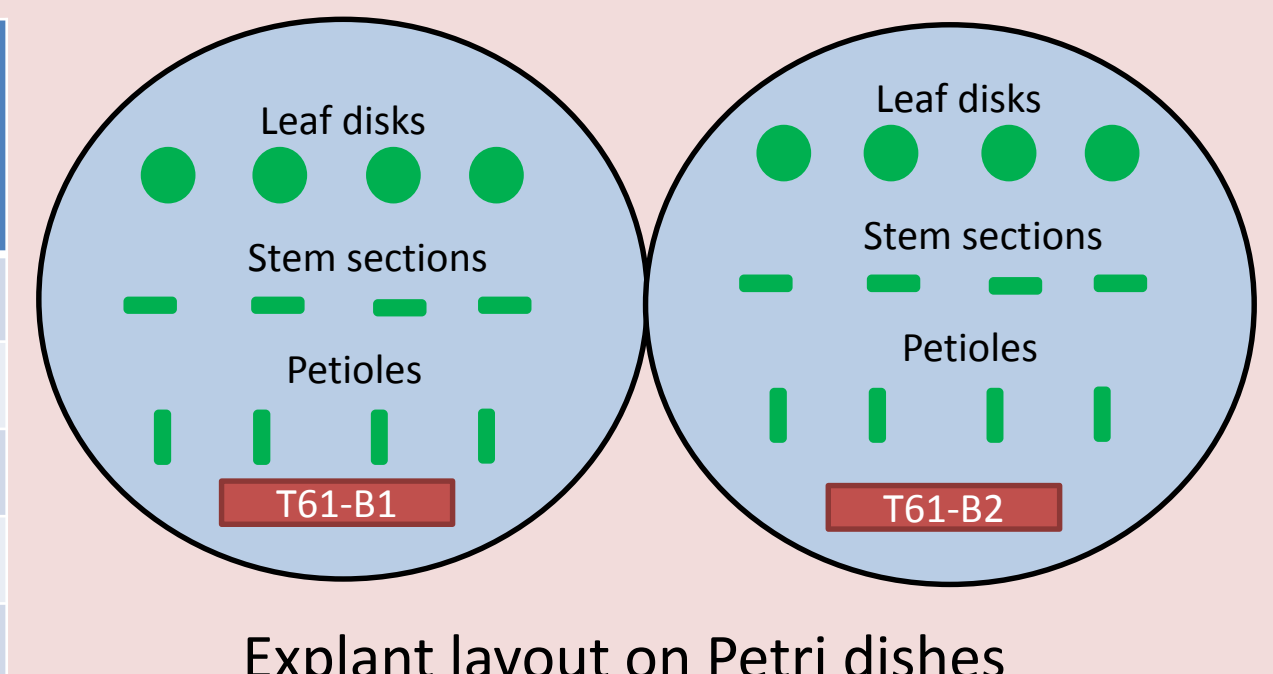


We studied genetic variation in shoot regeneration in five basal media

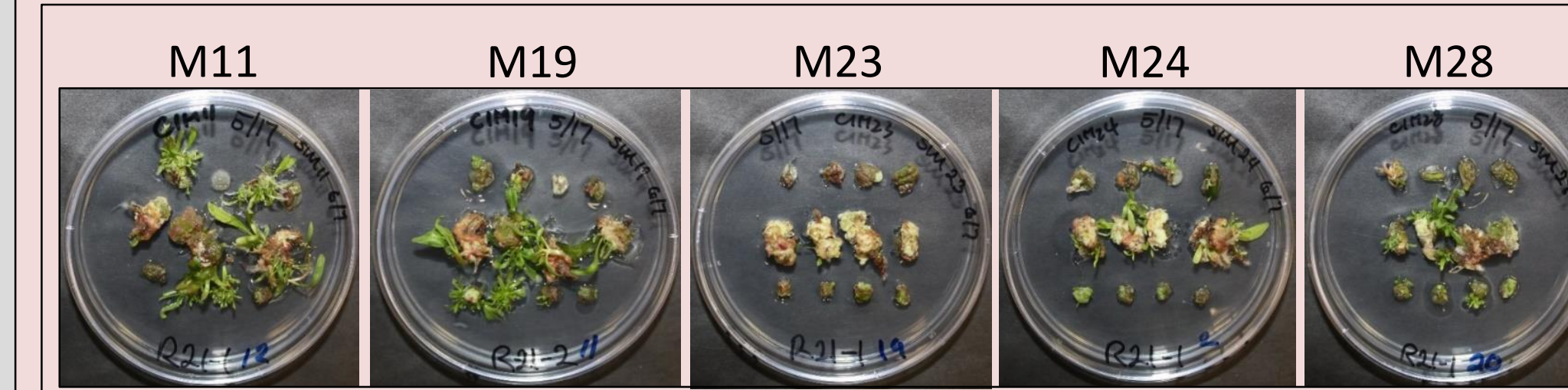
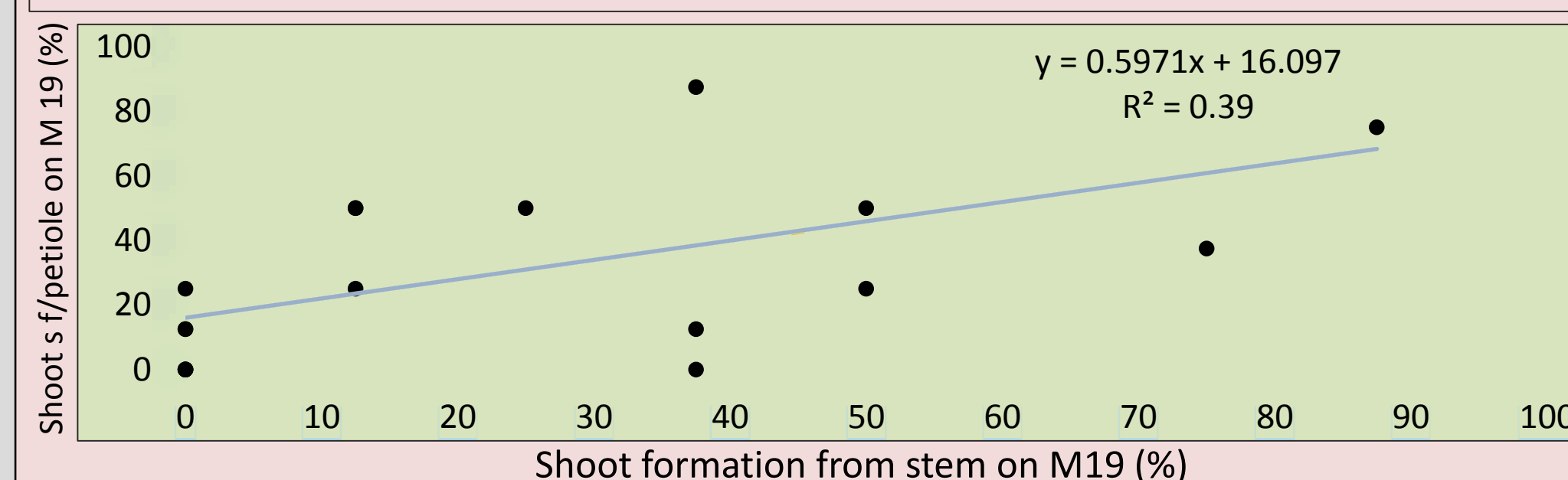
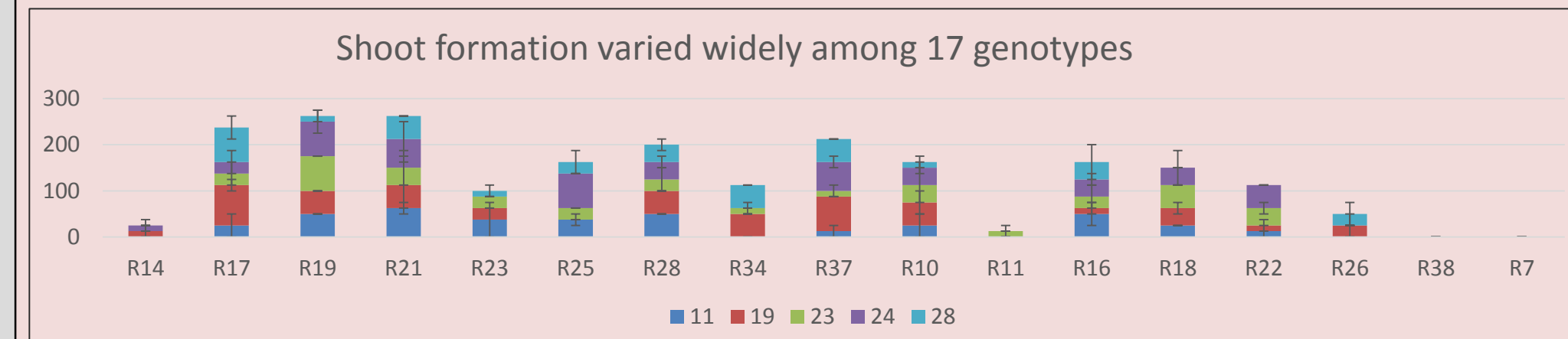
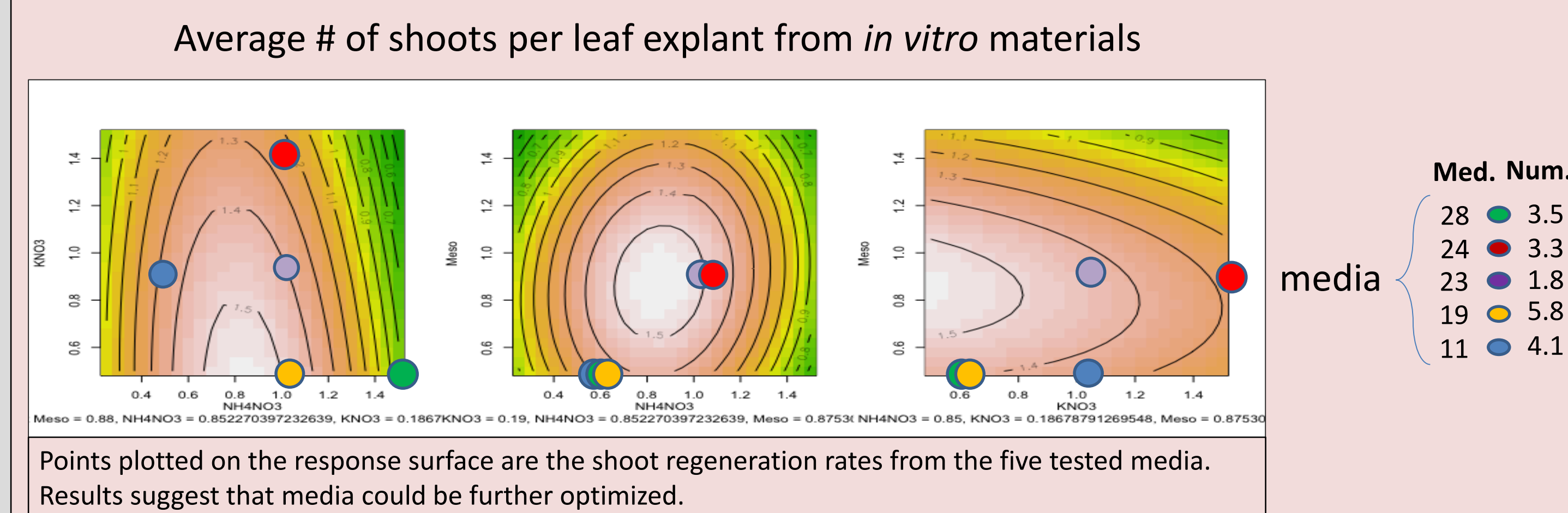
Five basal media that gave good regeneration were tested in an indirect regeneration system (Table 2). One experiment had leaf, stem and petiole explants derived from 6 genotypes of *in vitro* grown plants, and a second expt. had the same explant types from 17 genotypes of *in vivo* grown plants.

Table 2. Nutrient components of tested media.

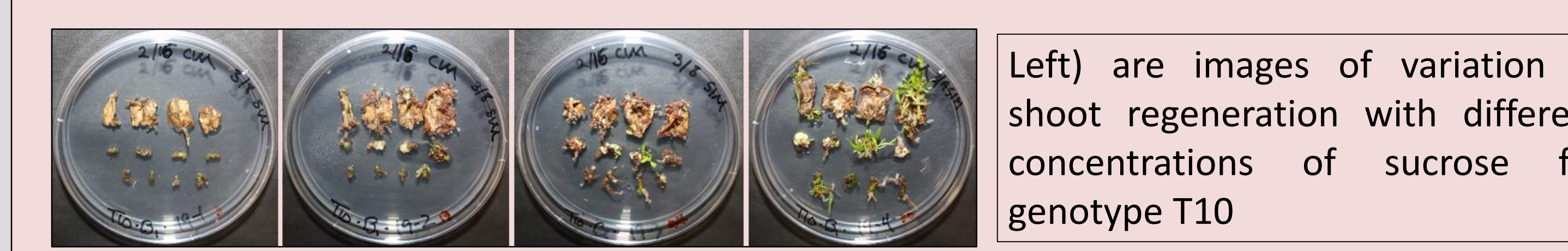
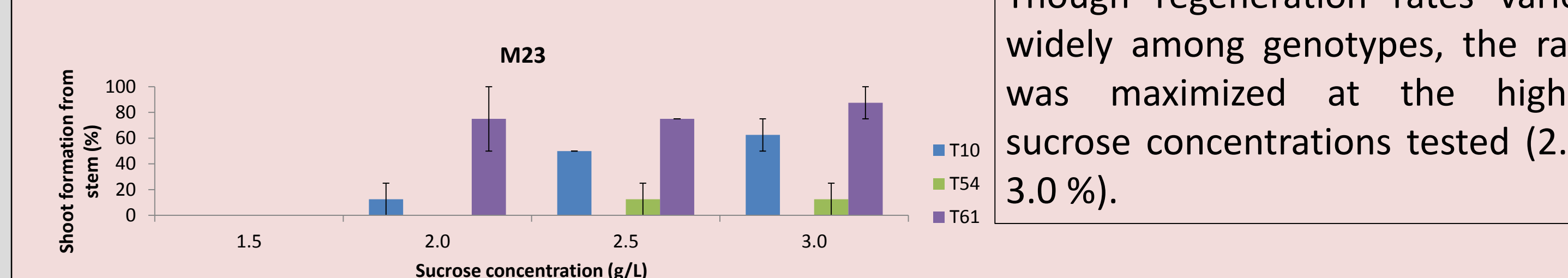
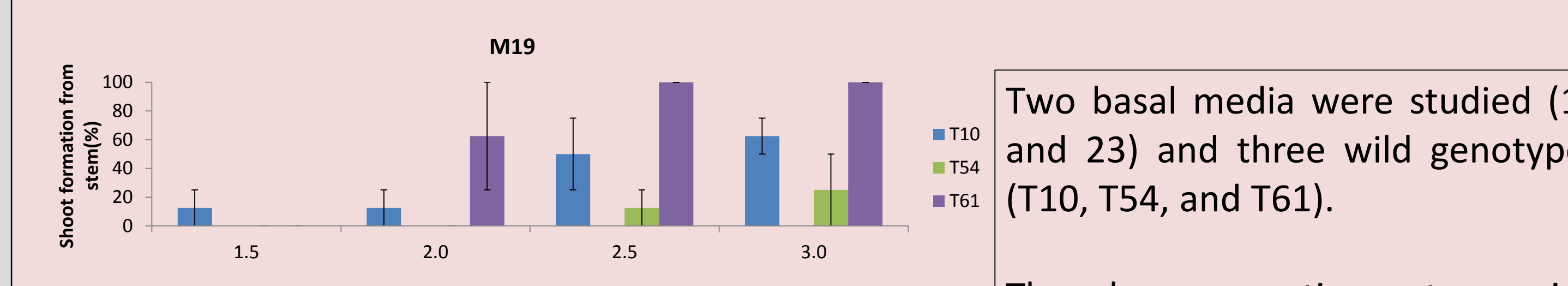
Treatment	NH ₄ NO ₃	KNO ₃	Mesos: CaCl ₂ , MgSO ₄ , KH ₂ PO ₄	Micro-nutrients	Fe-EDTA
11	0.5X	1X	0.5X	1X	1X
19	1X	0.5X	0.5X	1X	1X
23	1X	1X	1X	1X	1X
24	1X	1.5X	1X	1X	1X
28	1.5X	0.5X	0.5X	1X	1X



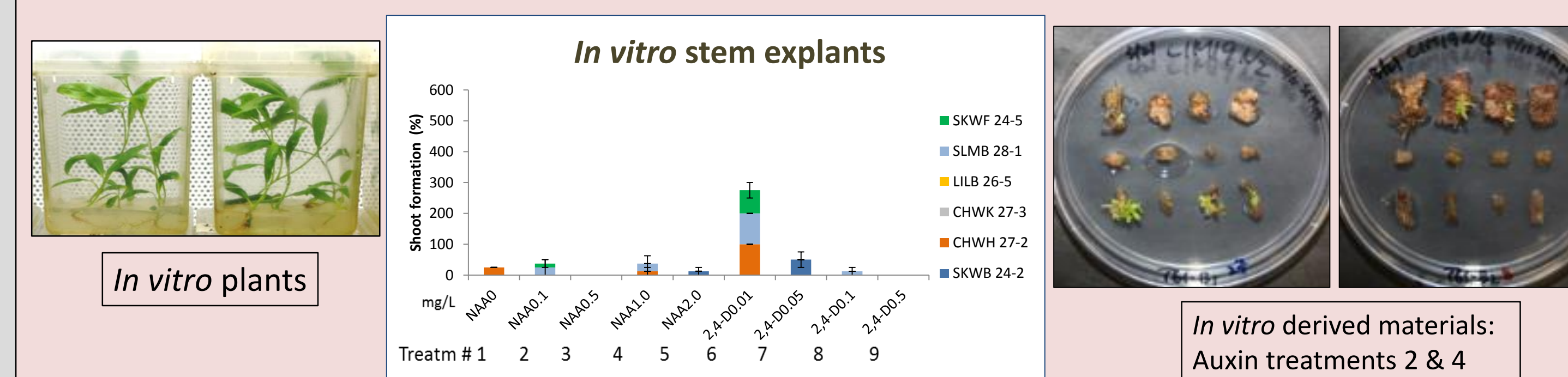
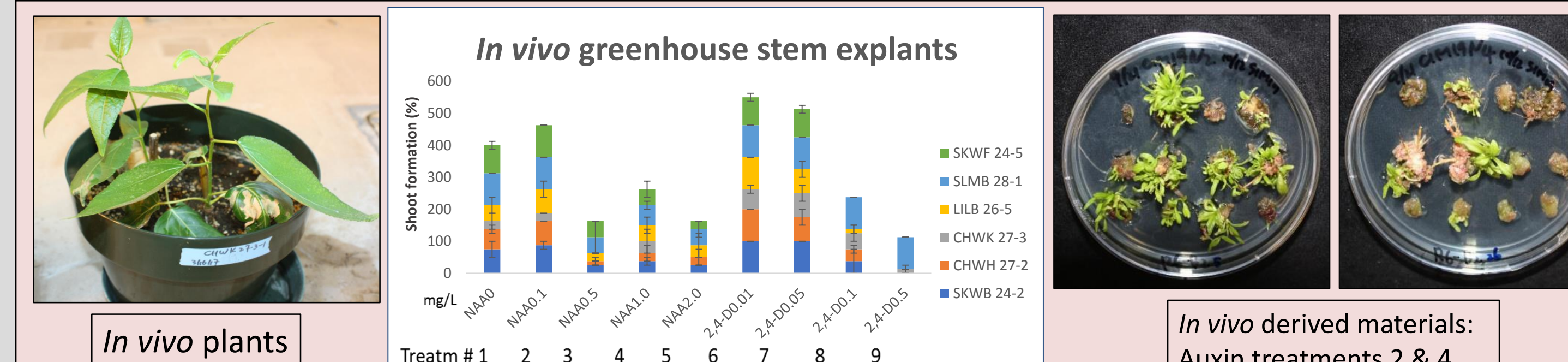
Medium 19 gave highest shoot regeneration from leaf & stem explants



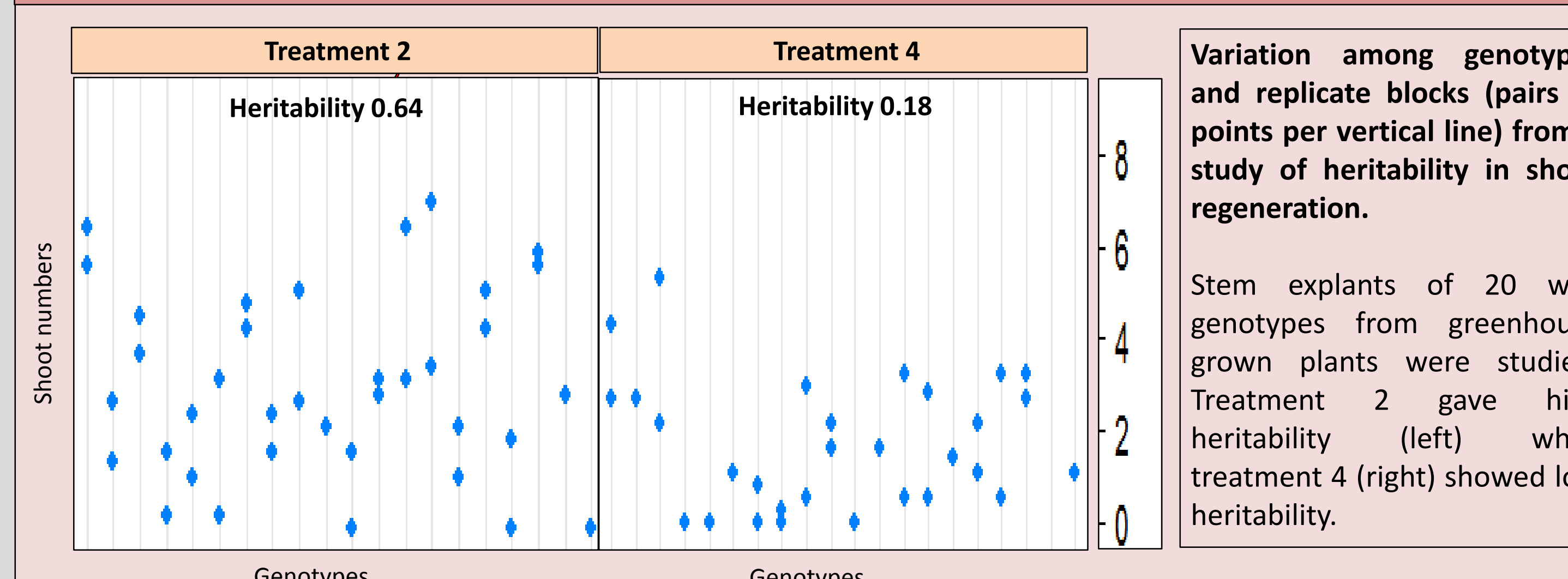
High concentrations of sucrose enhanced shoot formation from in vitro stems



Explants from in vivo greenhouse plants had superior shoot formation to explants from in vitro plants

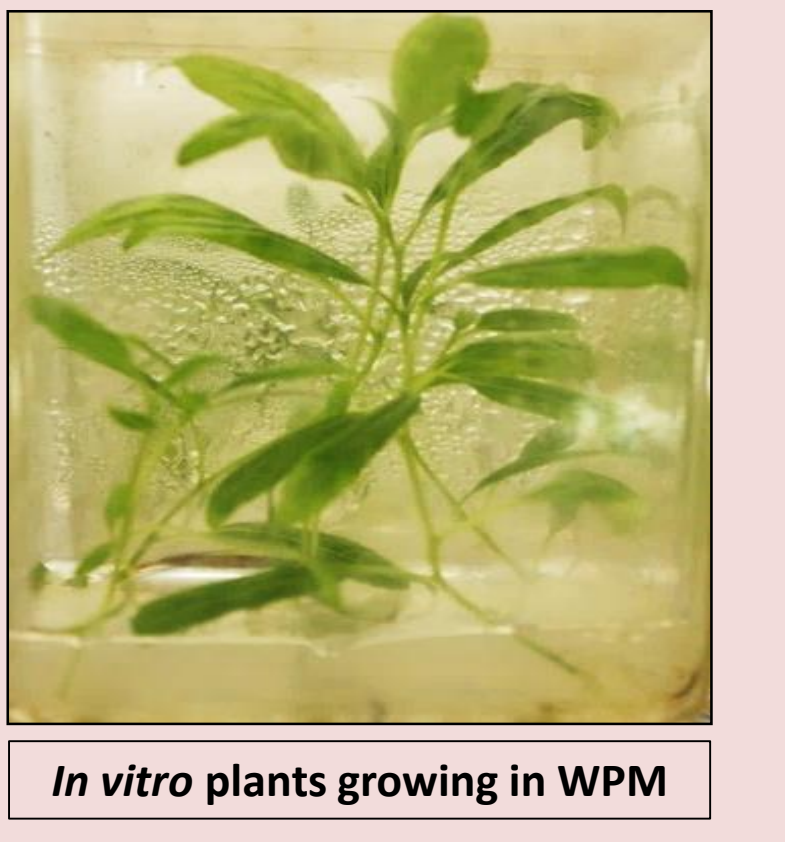


Broad-sense heritability varied widely among different auxin treatments, and is a useful parameter for helping to select optimal treatments for use in GWAS studies

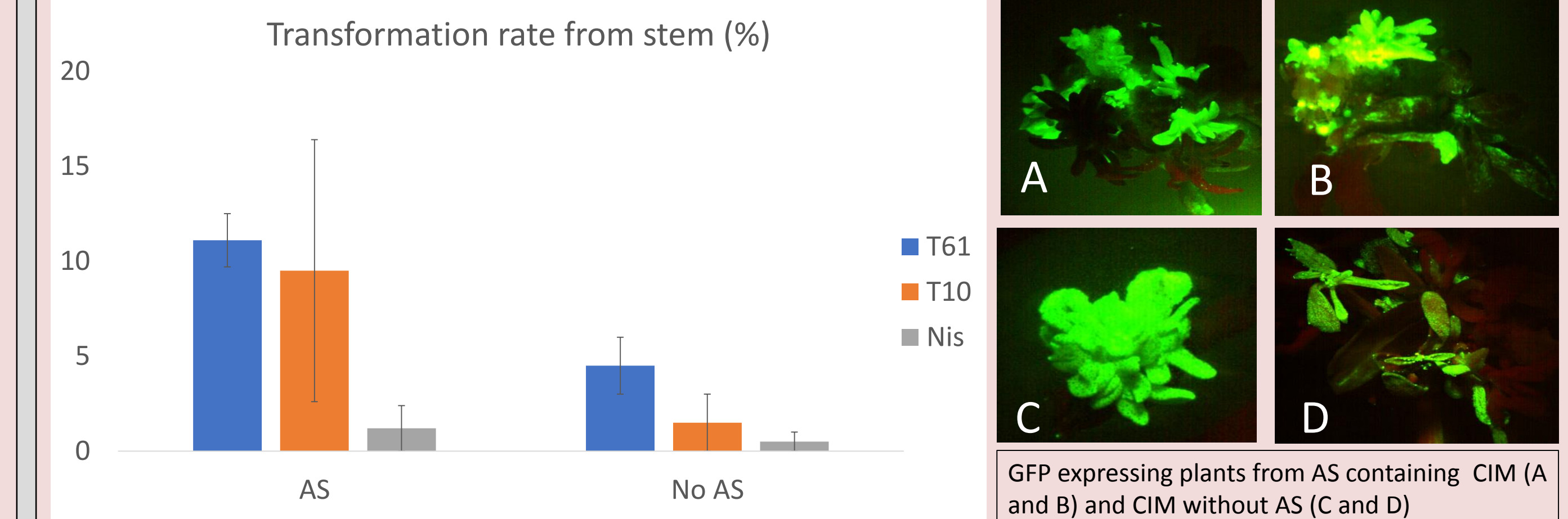


Transformation protocol development for Populus trichocarpa

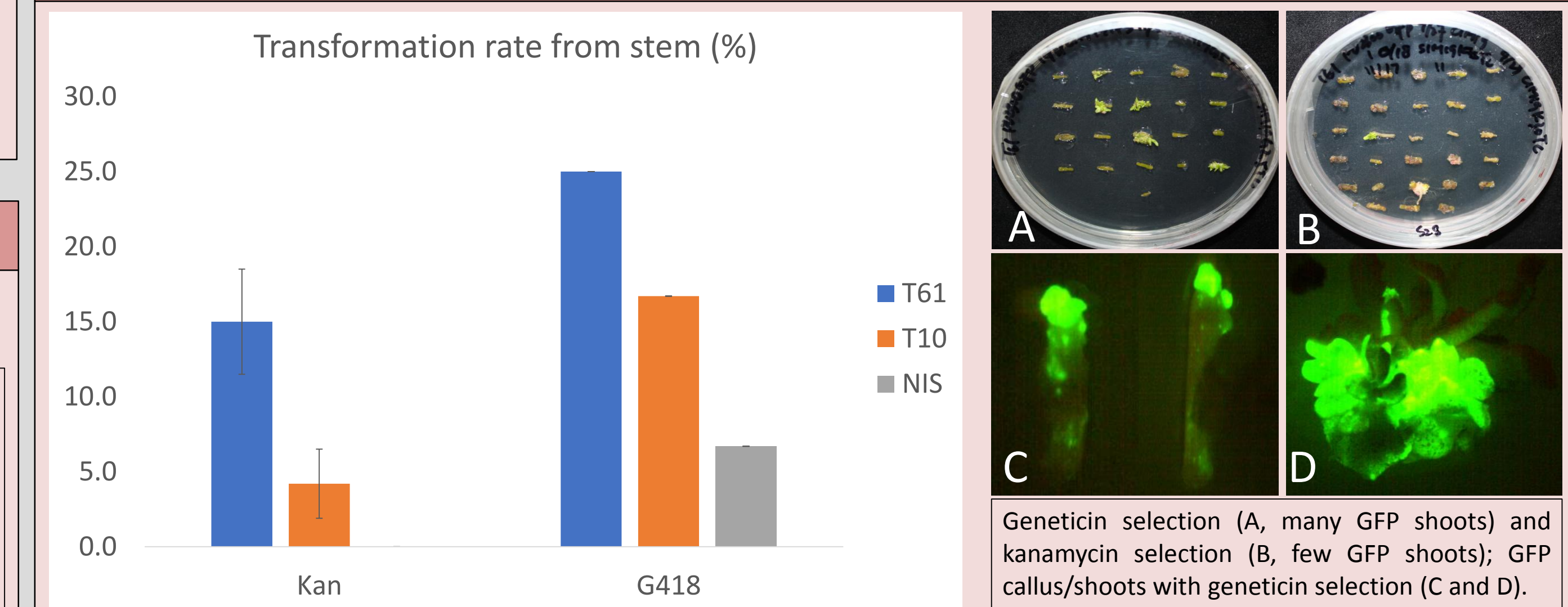
- Three genotypes were randomly selected for study
- Plants were grown in WPM hormone-free medium (example of source plants shown to right)
- Leaf and stem (including petiole) explants were co-cultivated with 2x35S::eGFP in Pc2300 (contains NPTII selection)
- Two to four plates per genotype and 20-30 explants per plate
- Stable GFP expression verified under GFP microscope
- Three factors studied were:
 - AS vs. no AS in co-cultivation CIM
 - Geneticin (G418) vs. kanamycin selection
 - Duration on CIM with G418 selection



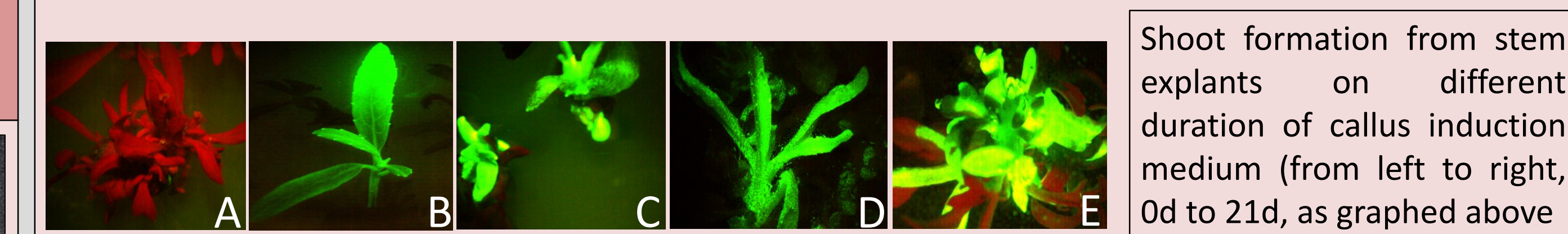
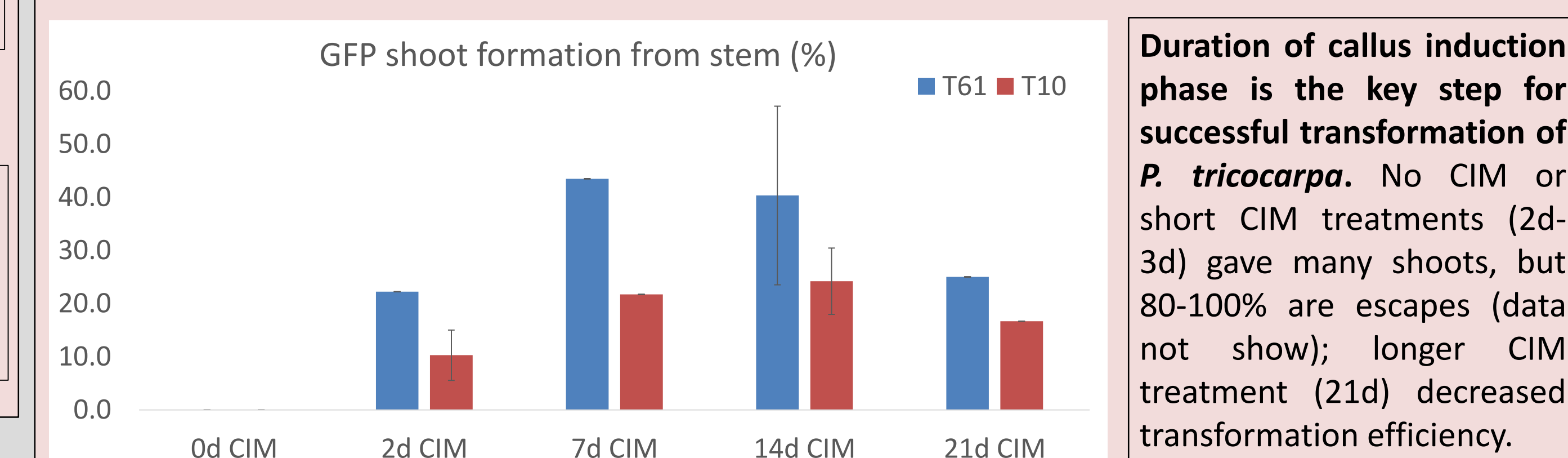
Transformation efficiency doubled after 2 days of co-cultivation on CIM containing 50 μM AS



Geneticin (5 mg/L) selection increased transformation rates over that from kanamycin selection for three genotypes tested



Transformation efficiency was doubled in genotypes T10 and T61 when CIM was optimized at 7 to 14 days for each genotype



Conclusions

- Shoot regeneration from two tested genotypes of *P. trichocarpa* varied widely among different strengths of macro and micro element-containing media of 64 tested
- Among 5 selected basal media, low KNO₃ and MESO (M19) gave better organogenesis for all genotypes; this basal medium was selected for further optimization treatments
- Explants from greenhouse-grown *in vivo* plants showed markedly higher shoot regeneration than from *in vitro* grown plants
- Shoot regeneration was greatest at a high level of sucrose for all genotypes
- Acetosyringone induction in CIM during co-cultivation with *Agrobacterium* greatly improved the rate and intensity of stable GFP from stem explants
- Geneticin was far more efficient for selection and regeneration of transgenic callus and shoot selection than was kanamycin
- 7 days and 14 days on CIM with selection greatly improved transgenic shoot recovery compared to our standard CIM treatment of 21 days
- Auxin concentrations and types affect rate and heritability of shoot regeneration, and are thus important to consider when selecting conditions for GWAS

Acknowledgements

We thank the National Science Foundation Plant Genome Research Program (IOS # 1546900) and the Tree Biosafety and Genomics Research Cooperative at Oregon State University for support

