

# Screening of Cultivars for Tissue Culture Response and Establishment of Genetic Transformation in a High-yielding and Disease-resistant Cultivar of *Theobroma cacao*

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## Abstract

A highly efficient transformation protocol is a prerequisite to developing genetically modified and genome-edited crops. A tissue culture system spanning the initiation of floral material to the regeneration of plantlets into soil has been tested and improved in cacao. Fourteen cultivars were screened for their tissue culture response and transfer DNA (T-DNA) delivery efficiency via *Agrobacterium*. These key factors were used to determine the genetic transformability of various cultivars. The high-yielding, disease-resistant cultivar INIAPG-038 was selected for stable transformation and the method was further optimized. Multiple transgenic events were produced using two vectors containing both yellow fluorescent protein and neomycin phosphotransferase II genes. A two-fold strategy to improve both T-DNA delivery and secondary somatic embryogenesis rates was conducted to improve overall transformation frequency. The use of *Agrobacterium* strain AGL1 and cotyledon tissue derived from immature somatic embryos ranging in size between 4-10 mm resulted in the highest T-DNA delivery efficiency. Furthermore, the use of higher concentrations of basal salts and cupric sulfate in secondary callus growth medium increased the percentage of explants producing greater than ten embryos by 504% and 443%, respectively. Consequently, an optimal combination of all these components resulted in a successful transformation of INIAPG-038 with 3.7% frequency at the T<sub>0</sub> plant level. Grafting transgenic scions with undeveloped roots to wild-type seedlings with strong, healthy roots helped make plantlets survive and facilitated quick transplantation to the soil. The present methods can be applied to improve tissue culture response and transformation frequency in other cacao cultivars.

## Key message

Tissue culture and genetic transformation methods for a high-yielding, disease-resistant cultivar of *Theobroma cacao* were established while factors affecting T-DNA delivery and somatic embryogenesis were identified.

**Keywords** *Theobroma cacao* – *Agrobacterium tumefaciens* – Genetic transformation – Somatic embryogenesis – Regeneration – Grafting

## **Abbreviations**

**DTT:** Dithiothreitol

**ED3:** Embryo development 3% sucrose

**ED4:** Embryo development 4% sucrose

**ED6:** Embryo development 6% sucrose

**EDL:** Embryo development light

**FPE:** Fluorescent protein expression

**nptII:** Neomycin phosphotranferase II

**OD600:** Optical density at 600nm

**PCG:** Primary callus growth

**QSE:** Quaternary somatic embryo

**SCG:** Secondary callus growth

**T-DNA:** Transfer-DNA

**TSE:** Tertiary somatic embryo

**YFP:** Yellow fluorescent protein

## **Introduction**

*Theobroma cacao* L. (cacao) is an economically important crop grown predominantly in Africa, although the species was originally domesticated in South America and significant production still remains there. Despite cacao beans being a commodity in a multibillion-dollar industry, there have been severe losses from disease to both yields and trees (Fister et al. 2018; Marelli et al. 2019). Obstacles such as long juvenile period, issues with heterozygous and heterogeneous populations (Wickramasuriya and Dunwell 2018),

long-life cycle and funding constraints on conventional cacao breeding programs (Maximova et al. 2003) make desired traits like high yield, disease resistance and bean quality difficult to combine into a single cultivar. However, both genome editing and transgenic approaches can be used to improve and greatly accelerate the pace of trait development.

The first stable transformation to regenerate cacao plants was reported by Maximova et al. (2003) and since then there have been other successful reports of the transformation of TcChil1 (Maximova et al. 2006) and TcLEC2 (Shires et al. 2017) into cacao. The introduction of BABY BOOM transcription factor into cacao significantly increased transformation frequency, but it caused a regeneration issue with an abnormal phenotype (Florez et al. 2015). These previously mentioned experiments were performed on the amenable cultivar PSU SCA-6. Previous to this study, PSU SCA-6 was the only cultivar that transgenic plants have been regenerated from. Although PSU SCA-6 is not identical to SCA-6, the original SCA-6 was collected from the wild in the Peruvian Amazon (Zhang et al. 2011) and has some natural, broad-spectrum fungal resistance (Pokou et al. 2019). However, this cultivar is not highly productive (Wahyu et al. 2009). For genetic transformation to be used most effectively the ability to transform many cacao cultivars, including those less-amenable to transformation, must be established. Ideally, a single genotype-independent transformation method would be developed, but similar to somatic embryogenesis which has had broad success across the plant kingdom, there still exists a high degree of genotype-to-genotype variation and so protocol customization is necessary (Garcia et al. 2016). Optimization of the basal salts, duration of hormone treatments, explant quality, and explant type are effective ways to improve the culturing response of recalcitrant cultivars.

Several explant types are used in cacao tissue culture and the transformation process, which include petals and staminodes from immature flower pods, cotyledon tissue from immature somatic embryos, and mature somatic embryos. Petals and staminodes are capable of primary somatic embryogenesis (Maximova et al. 2002) and are the starting point for all subsequent processes. Cotyledon tissue derived from somatic embryos is an alternative tissue type, efficient at secondary, tertiary and quaternary somatic embryogenic processes (Maximova et al. 2003). Mature somatic embryos undergo germination and are able to regenerate



































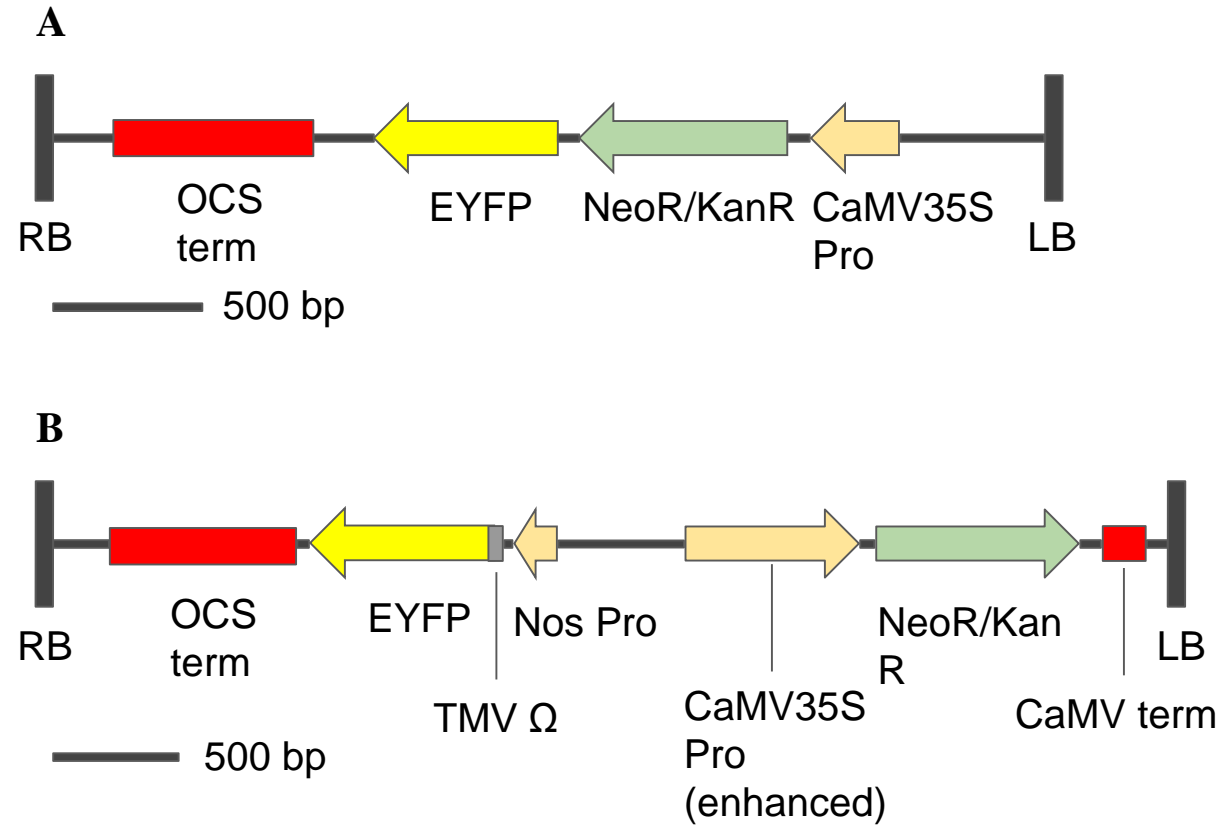












**Fig. 1** Schematic diagram of two transformation vectors used for cacao transformation. **(a)** pDDNPTYFP-1 is a 3637-bp T-DNA fragment containing *nptII* and *eyfp* translational fusion driven by a CaMV35S promoter. **(b)** pDDNPTYFP-2 is a 4416-bp T-DNA fragment containing a two gene concept where one gene has *eyfp* driven by a Nos promoter/TMV Ω enhancer and the other gene has *nptII* driven by an enhanced CaMV35S promoter























