

Maize Transformation with Feedback- Insensitive Anthranilate Synthase Gene by Particle Bombardment of Immature Embryo

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ABSTRACT

Anthranilate synthase (AS) is a key enzyme in the synthesis of tryptophan (Trp). Tryptophan is one of the essential amino acids and the major contributor of the indole ring for the synthesis of many important compounds including auxins, glucosinolates, nicotinic acid, phytoalexins and alkaloids. Anthranilate synthase (AS) catalyzes the first reaction in the multi-step Trp biosynthesis pathway by converting chorismate to anthranilate. Particle bombardment is a popular method currently used for producing transgenic maize. Particle bombardment was carried out using a plasmid pC2ASA2-NOS-ASB with (feedback anthranilate synthase) ASA2 (α & β) genes under control of CaMV35S promoter. The best conditions to transform maize variety Hi_{II} were obtained when immature embryos were cultivated, prior to the bombardment, in higher osmolarity for 4 hours and bombarded at an acceleration helium gas pressure of 900 and 1,100 psi single and double shot. (6methyl – D-L- tryptophan) 6 MT 100 μ M used on selection medium as selective agent. PCR for nptII, ASA2, ASB genes and Southern blot using ASA2 probe were used to detect insertion of gene. The results clarified that two lines showed 12-16 fold increase in the tryptophan level compared with wild type.

Keywords : nuclear transformation, immature embryo culture, 6methyl-D-L- tryptophan, plant regeneration , transformation.

INTRODUCTION

Maize is a major commodity in international agriculture and an important source of protein and energy for human and livestock nutrition; it has been one of the prime targets for genetic manipulation. However, most studies on maize transformation have utilized genotypes adapted to temperate zone. Maize (*Zea mays* L.) is the third most planted cereal crop after wheat and rice worldwide. However, the production of this crop is decreasing due to increased population, limited land, environmental and biotic stresses over the years. Conventional breeding has been used as a tool to overcome these constraints, this has resulted in development of modest increments in yields and agronomic characteristics such as disease resistance and drought adaptability to different agroecological zones (Ayaga, 2003).

Biolistic transformation of maize has become routine since the recovery of fertile transgenic plants by this method was first reported by Gordon-Kamm et al. (1990). This technology has been used to transform various maize target tissues (Gordon-Kamm et al., 1999), including immature zygotic embryos from inbred lines (Koziel et al., 1993; Dunder et al., 1995; Brettschneider et al., 1997) and Hi II germplasm (Songstad et al., 1996; Pareddy et al., 1997). Immature zygotic embryos provide an excellent transformation target because they minimize both time in tissue culture and the expertise needed for callus induction and maintenance (Brettschneider et al., 1997; Songstad et al., 1996).

The regeneration of maize in tissue culture is important for the production of transgenic maize and for crop improvement using genetic engineering approaches. The first somatic embryos in maize tissue culture were produced by (Green and Phillips 1975) The enormous progress achieved by molecular and cellular biology in recent years has generated a new understanding for many basic genetic concepts and biochemical mechanisms, allowing the emergence of improved strategies for breeding and cultivar development. Among the new generated technologies, plant transformation is an important tool for the genetic manipulation and improvement of crop species (Carneiro et al., 2000). Molecular breeding methods involving marker assisted selection and genetic transformation now provide viable alternatives in several crops, including maize (Frame et al., 2002). However, the pre-requisite for crop genetic transformation is existence of a reliable plant regeneration system.

To transfer foreign genes into the maize genome, various reports have shown that microprojectile bombardment is a successful technique (Klein et al., 1988; Brettschneider et al., 1997; Frame et al., 2000; Lorence and Verpoorte 2004; Herrera-Estrella et al., 2005).

Plant transgenic technology has undergone key changes in the past 30 years; major technological developments have been sustained by breakthrough studies by investigators all over the world. However, there are still a number of bottlenecks in the technology used to transform plant species. (Vain 2007). Tryptophan (Trp) is one of the essential amino acids and the major contributor of the indole ring for the synthesis of many important compounds including auxins, glucosinolates, nicotinic acid, phytoalexins and alkaloids (Haslam 1993). Trp is synthesized by a five-enzyme biosynthetic pathway from the branch point compound chorismic acid at the end of the shikimic acid pathway (Radwanski and Last 1995). The first step of this five-enzyme pathway is catalyzed by anthranilate

synthase (AS), which converts chorismate to anthranilate. As a branch point enzyme in aromatic amino acid biosynthesis AS has an important regulatory role through feedback inhibition by Trp (Widholm 1974; Singh and Widholm 1974). Plant AS is a heterotetramer of two α - and two β -subunits encoded by separate nuclear genes and synthesized in the cytosol, as precursor proteins that are then localized in the plastids (Poulsen et al. 1993; Romero et al. 1995). The α -subunit binds the substrate chorismate and carries out its aromatization while the β -subunit transfers an amino group from the other substrate glutamine (Crawford 1989). This glutamine-dependent AS reaction requires both α - and β -subunits. The α -subunit alone can synthesize anthranilate from chorismate using ammonia as the amino donor rather than glutamine, if ammonia is present in high concentrations. The AS reaction is normally subjected to feedback inhibition by Trp which binds to an allosteric site on the AS α -subunit (Bohlmann et al. 1996; Li and Last 1996). In addition to Trp, different Trp and indole analogs that can be converted to Trp analogs (Widholm 1981) can also inhibit AS and in turn plant cell growth.

The final objective of this study is to get transformed plants with feedback-anthranilate synthase and used 6-methyl-DL-tryptophan (6MT) as selection agent instead of most commonly used 5-methyltryptophan (5MT). The maize genetic transformation protocol developed in this work will possibly improve the efficiency to produce new transgenic maize lines expressing desirable biochemical pathway.

MATERIALS AND METHODS

1- Preparation of the culture

Immature embryos, between 1 and 2 mm length (14-16 days after pollination) were harvested from plants grown in the green house, Illinois University. Ears for embryo extraction were surface-sterilized using 70% alcohol for 5 min then 10% sodium hypochlorite was used for 20 min and rinsed three-five times with sterilized distilled water. Immature embryos were isolated and cultivated scutellum-side up onto the surface of D medium (Duncan et al., 1985) for 21 days then to N6 salts and N6 vitamins (Chu et al., 1975), 2 mg/l 2,4-D, 3% sucrose, 100 mg/L myo-inositol, 2.76 g/L proline, 100 mg/L casein hydrolysate and 2.5% gelrite, pH 5.8, filter sterilized silver nitrate (25 μ M) added after autoclaving. Cultures were kept in Petri dishes at 28°C and grown in the dark for callus initiation and maintenance.

2- plasmid

The plasmid used in this work, pC2ASA2-NOS-ASB 16.6 kb (Illinois University), (fig, 1) contains a selectable marker (ASA2 α & β) the coding region of kanamycin, *gusA*, under control of the cauliflower mosaic virus (CAMV). ANOS polyA (nopaline synthase) terminator sequence

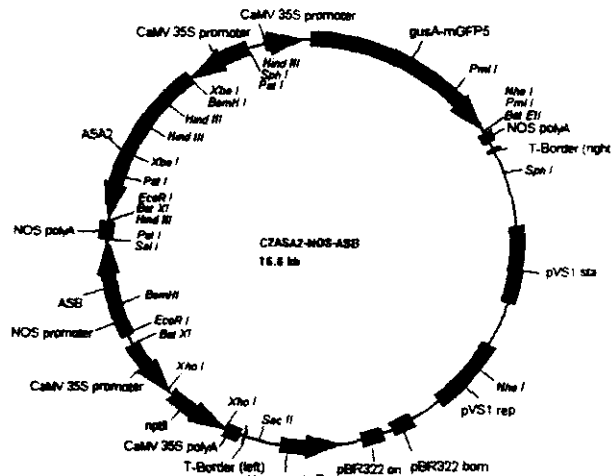


Figure 1- Schematic drawing of the plasmid C2ASA2-NOS-ASB. The plasmid 16.6 kb consists of ASA2, ASB, CaMV35S : cauliflower mosaic virus 35 S promoter and 3'NOS : the polyadenylation signal of nopaline synthase.

3- Maize transformation and regeneration

Friable callus was used for particle bombardment. Callus was transferred to osmoticum medium (N6 medium + 36.4 g/L sorbitol and 36.4 g/L mannitol) for 4 hours prior to bombardment. The gold particles (Bio-Rad) (1 μ g) was used to precipitate DNA onto the microparticles 1 μ L plasmid DNA (stock 1 μ g μ L⁻¹). Then, 220 μ L (stock 2.5 M) and 50 μ L spermidine (stock 0.1M) were added and homogenized. The mixture was kept on ice for 5 min and vortexed for 5 min, micro centrifuged at 5000 rpm for 1 min, rinsed carefully with 250 μ l of ethanol and suspended in 40 μ L 100% ethanol. 10 μ l of the DNA -coated particles were pipetted onto each macrocarrier (washed in absolute EtOH, dried before uses). Bombardments

were performed on Petri dishes containing friable callus clump in the middle. Different treatments were designed to test the pressure of the accelerating helium pulse (900 and 1100 psi single and double shout). Selection of putative transformed callus began 10 days after bombardment when the callus were cultured every 21 days on N6S selection medium which was similar to callus initiation medium but without proline and casein hydrolysate, filter sterilized silver nitrate (5 μ M) was added after autoclaving, 6- Methyl-DL-tryptophan (6MT) 100 μ M was added also.

The surviving callus was transferred to regeneration medium MS medium (Murashige and Skoog 1962) supplemented with 1ml/L (1000X) MS vitamin stock, 100mg/L myo-inositol, 60 g/Sucrose , 3g/L gelrite, pH5.8 ,after autoclaving 100 mg/mL of 6MT was added. The calluses were incubated for 2 weeks at 25°C in the dark. After 2 weeks, transformed green calluses were transferred to the light on Regeneration medium (II) which is the same for MS I except MS II supplemented with 30 g/L sucrose. For the reporter gene *gusA* transient expression studies, 20% of bombardment callus was incubated two days after bombardment in a solution containing 380 μ l 1M Na₂HPO₄, 620 μ l 1M NaH₂PO₄, 200 μ l 0.5 mM EDTA, 1000 μ l 0.5 mM K-ferricyanide, 1000 μ l 0.5 mM K-ferrocyanide, X-Glue (10 mg in 40 μ l of DMF) and 6760 μ l water. Vacuum for 5-10 min, avoid air bubbles and floating samples, incubate on 37°C over night, sealed with parafilm. Gus spots appeared.

4- PCR analysis and Southern blotting

For the molecular characterization of the transgenic plants generated, total genomic DNA was isolated from leaf tissue of primary leaves using a CTAB protocol described by (Doyle and Doyle 1990) and submitted to PCR and southern blot analysis. For the presence of ASB, the primers 5' TGTCCAAGATCCCATGACGATTCC 3' and 5' CAGAAATCCACAGAACCGGGAGAT 3', which amplify 800 bp were used. For the ASA2 (α & β), the primers 5'TCTGTACACTTCAAATGGGTCAGC3' and 5' CTAAAAGCGGGAACCTTGATTCCGC3' which amplify 815 bp were used and for *nptII* the primers 5'ATCTCACCTTGCTCCTGC3' and 5'ATACCGTAAAGCAGGAGG3' which amplify 200bp were used. Each 20 μ l amplification reactions mixture consisted of 100ng of template DNA, 2 μ l of Taq buffer 10X, 0.5 μ l of dNTPS 10 mM, 1 μ l of each primer, and 0.3 μ l of Taq polymerase. The reactions were carried out using a thermal conditions: 1 cycle at 94°C for 5 min, 35 cycles at 94°C for 45s, annealing at 54.6°C for 45s , 72°C for 60s and a final extension at 72°C for 7 min for GUS, 1 cycle at 94°C for 5 min, 35 cycles at 94°C for 30s, annealing at 57°C for 45s

,72°C for 60s and a final extension at 72°C for 7 min for ASA2, and 1 cycle at 94°C for 5 min, 30 cycles at 94°C for 45s, annealing at 55°C for 30s, 72°C for 45s and a final extension at 72°C for 7 min. The amplified products were separated by electrophoresis on 1% agarose gel and visualized with ethidium bromide stain under UV light.

For southern blot analysis, 15 micrograms of total genomic DNA from each T0 plant were completely digested with BamH1 at 37°C overnight, separated by electrophoresis in 1% agarose gel, and then transferred onto a Hybond- N⁺ nylon membrane according to (Sambrook et al., 1989). The blot was hybridized with a ³²P labeled ASA2 gene coding region. Hybridized membranes were exposed to Kodak film and kept at -80°C for 7 days.

The youngest expanded leaf of a shoot culture was ground into coarse powder in liquid nitrogen. The leaves powders were homogenized in 5 volumes of 0.1 N HCl by the fast prep FP120 Cell Disrupter (Savant Instruments, Holbrook, N.Y.) at 4.0 ms⁻¹ for 40 sec. The supernatant was transferred to a new tube after centrifugation at 4°C and 13,000 g for 10 min. Another five volumes of 0.1 N HCl were added to the pellet, followed by homogenation and centrifugation as described above. The procedure was repeated 1 more time and the resultant were 15 volumes of supernatant was filtered with an Ultra Free-MC filter unit (Millipore, Bedford, Mass). Trp concentrations were measured by HPLC according to Cho et al. (2000).

5- Statistical analysis

Chi square test of homogeneity was used to determine the significance of differences.

RESULTS AND DISCUSSION

1- Effect of callus morphology on transformation efficiency:

The immature embryos of Hill were produced from selections of A188x B73, whose length was 1.6-2.0 mm. The embryos were selected and cultivated in D culture medium (Fig. 2a), then transferred to N6E culture medium. After 3 times of cultivation, two types of callus were produced. The present results are in accordance with those reported by (Vassil et al., 1991; Armstrong and Green, 1985) who found that Type I compact, white that failed to regenerate, Type II friable, yellowish and able to regenerate. Type II was selected and transferred for particle bombardment. (Figures 2b & 2c). When cultured *in vitro* can originate Type I or Type II embryogenic calluses. Type I callus is formed by hard, compact

and yellowish tissue, usually unable to regenerate plants. Type II callus is soft, friable, highly embryogenic and able to regenerate a higher number of plants than Type I callus. Type II embryogenic callus can regenerate only. On the other hand (Brettschneider et al., 1997; Frame et al., 2006), found that both types of calluses can be used to generate transgenic plants.

2- Effect of osmotic treatment on transformation efficiency:

An important aspect in transformation via biolistics is the damage to the target tissue during microparticle penetration into the cell. To minimize this problem, the target cells are usually plasmolysed by an osmotic treatment and increase stable clone recovery (Vain et al., 1993, Brettschneider et al., 1997) so callus were put on osmoticum medium for 4 hours before bombardment (Fig.2d). For bombardment, 900 and 1100 psi were used to shot calluses (single and double shot). The results indicated that double shot were more efficient than single shot. However, the differences between treatments were not significant according to Chi square test at $p < 0.05$ (0.3327). (Table 1) Clearing by demonstrates that the number of living calluses decreased with double shot but these living calluses showed positive results for PCR and are more efficient than single shot. (El-itriby et al., 2003) concluded from the transient GUS expression that, in the co-transformation experiments, the use of osmotic pre- and post-treatment with acceleration pressure of 1100 psi and double shots per plate is most efficient treatment.

3- Transient GUS activity

Gus expression was detected after 48 h in bombarded calluses. The blue spots appeared to confirm the insertion of the constructs (Fig. 2e). For selection medium, 6MT 100 μ M was used to select the transformed plants (Table 1, Fig 3a, 3b, 3c). (Abdel-Rahman and Widholm 2009) found that 6MT 100 μ M was the best concentration used for selection compared with 75 and 125 μ M. The use of feedback-insensitive AS α -subunit genes as selectable markers with 5MT as the selective agent has been reported in maize (Anderson et al., 1997) and rice (Tozawa et al., 2001).

4- Selection of transgenic maize plants over expressing ASA2

PCR analysis was conducted on all of the (133) plants regenerated from immature embryo transformation. It was observed 200 bp band of nptII gene (Fig.4a), 815bp band of ASA2gene (Fig. 4b) and 800 bp of ASB gene (Fig.4c). The result showed that (11) transgenic maize were positive for nptII, (57) were positive for ASB and ASA2gene. Southern blot of total

DNA from transgenic plants digested with BamHI and probed with ASA2 gene showed that feedback anthranilate synthase (high tryptophan) plants contained copies of the transgene (Fig 5) (Cho et al., 2004) showed that, the tobacco- feedback –insensitive ASA2 gene can be inserted into legume hairy roots (*A. sinicus* and soybean) and be expressed to produce feedback-insensitive AS activity that leads to increased free Trp levels and resistance to the Trp analog 5MT. The Trp analog 5MT and 6MT inhibit AS enzyme activity and plant growth. Over expression of the ASA2 gene results in resistance to 5MT in *E.coli* (Song et al., 1998) and transgenic *A. sinicus* (Cho et al., 2000).

The expression of ASA2 in plant tissues leads to increased levels of free Trp, which could be desirable since Trp. is an essential amino acid required in the diets of humans and non ruminant animals. To determine the over expression of the feedback- insensitive ASA2 had on Trp. production, free Trp levels in the T0 transgenic plant leaves were measured. The high- expresser lines 9 and 12 contained > 12-16 times, line 1 show the Trp. level as wild type (Fig 6). In this report, we demonstrate that a feedback-insensitive ASA2 gene can be used as a selectable marker for the production of transgenic maize plants. Since maize expressing ASA2 resistant to 6MT, thus ASA2 may be an effective selectable marker gene for use with many different species.

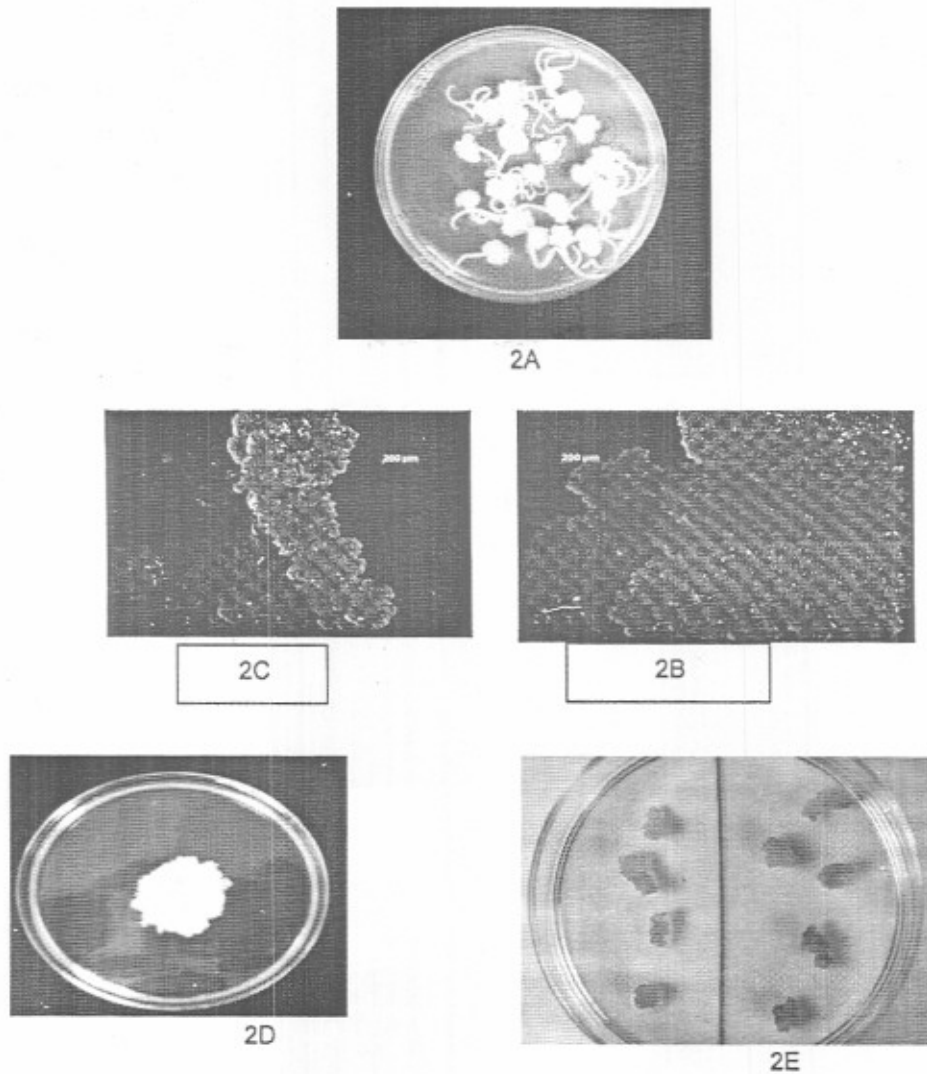
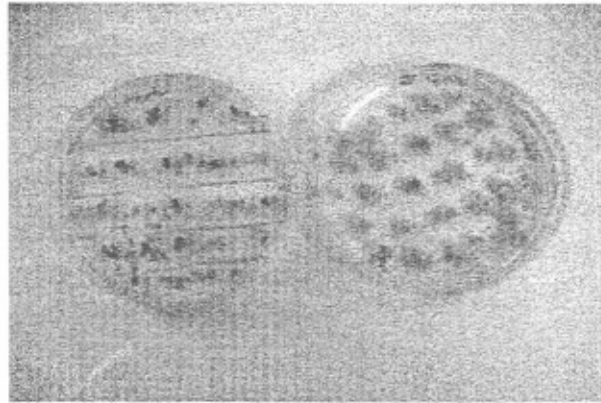
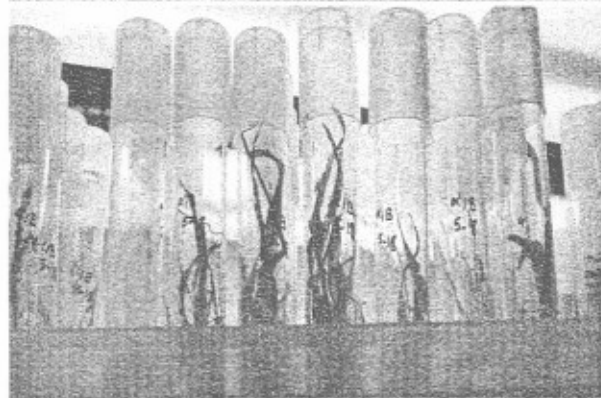


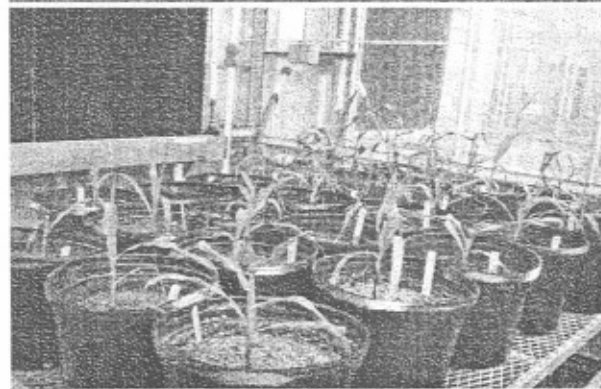
Figure (2) Morphology of different types of callus (A) immature embryo on medium D. (B) compact callus, (C) friable callus(embryogenic callus)(D) callus on osmotic medium before bombardment (E)transient GUS activity48 h after bombardment.



3A



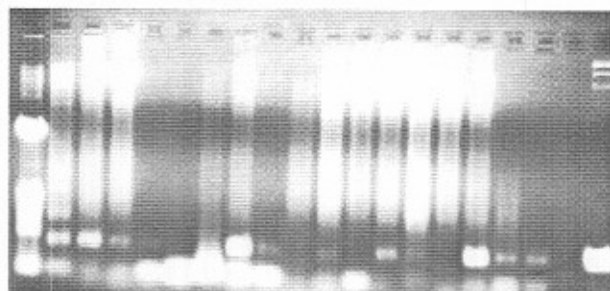
3B



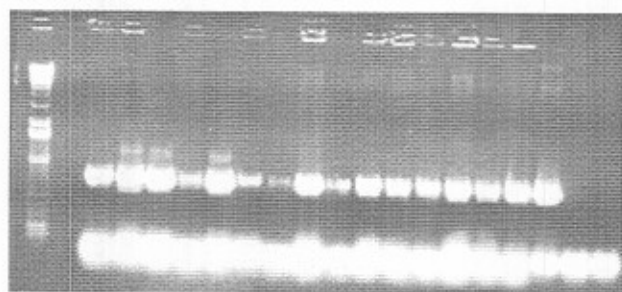
3C

Figure (3): Transgenic maize plants(A) calluses on selection medium with 6MT100 μ M,(B) shoot and root differentiation(C) Transgenic plants in the green house.

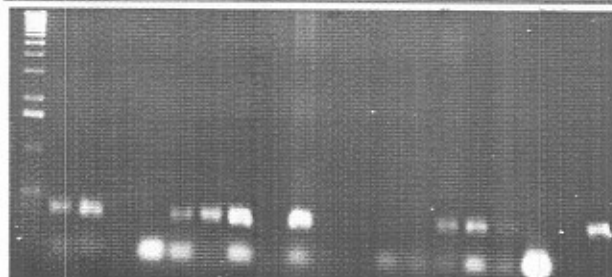
M, A₁, B₁, C₁, D₁, E₁, F₁, G₁, H₁, I₁, J₁, K₁, L₁, M₁, N₁, O₁, P₁, WT, W₁



4A



4B



4C

Figure (4) PCR analysis of regenerated plants. Agarose electrophoresis gels of PCR amplification products (A) co-transformed with nptII, (B) PCR products from ASA2 gene and (C) ASB gene.

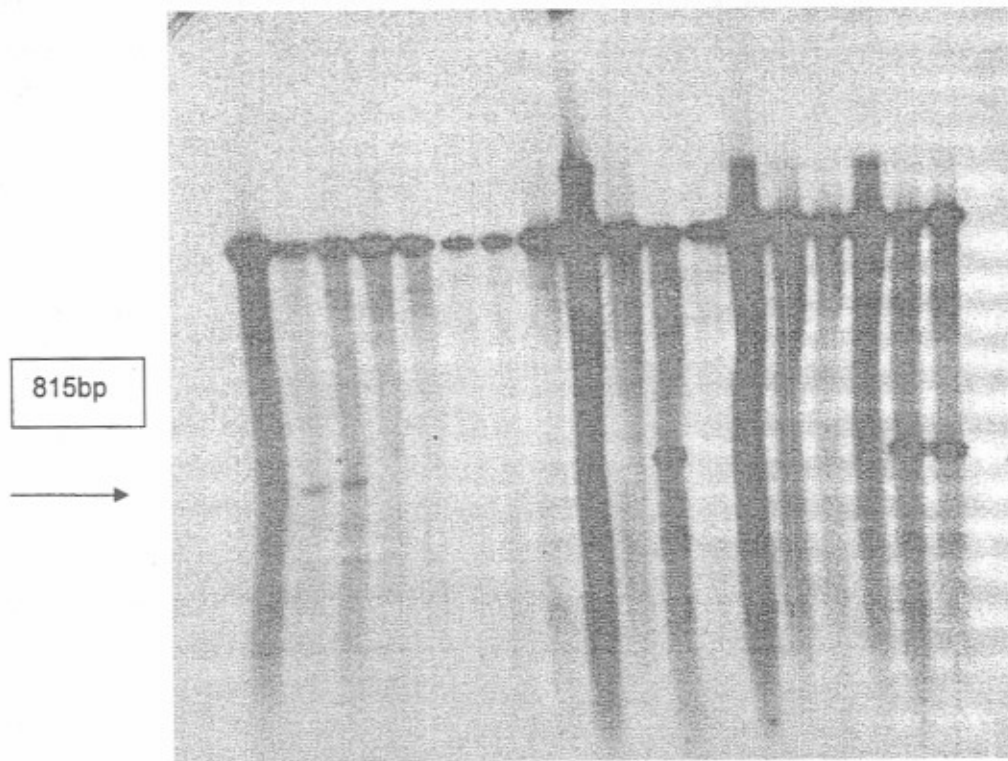


Figure (5): Southern analysis for ASA2 gene in transgenic plants

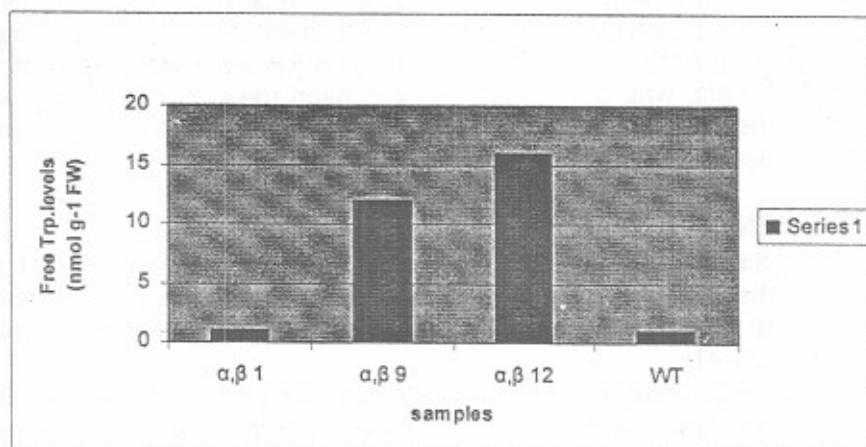


Figure (6): Free Trp levels in leaves of WT and T0 transgenic lines. Lines were selected on 100μM 6MT.

Table (1) # of calluses living and dead on selection medium N6S + 6MT 100μM

Helium acceleration pressure	Callus efficiency on selection medium						
	#of calluses	living calluses	% living calluses	Chi square	Dead calluses	% dead calluses	Chi square
single 900	49.67±12.93	7.84±4.71	16	0.054	41.72±15.78	84	0.00
single 1100	30.33±7.49	5.67±1.38	19	0.0145	24.56±5.25	81	0.00
double 900	20.34±3.94	3.0±0.84	15	0.0312	17.28±0.35	85	0.00
double 1100	20.67±8.38	4.0±1.34	19	0.206	16.74±2.81	81	0.00

Total chi square cal. = 0.3327 , df=3 p=0.05, Chi square tab. 7.82

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الملخص العربي

التحول الوراثي في الكالس الناتج من الجنين غير الناضج للذرة لإدخال جين التغذية الرجعية لتخليق إنزيم Anthranilate بواسطة قاذف الجينات.

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إنزيم ال Anthranilate synthase هو الإنزيم الرئيسي في تكوين الحمض الأميني التربتوفان. ويعتبر التربتوفان واحد من الأحماض الأمينية الرئيسية والمصدر الرئيسي لحلقة الإندول اللازمة لتخليق مركبات كثيرة ومهمة بما في ذلك الأوكسينات وحمض النيكوتينك والقلويدات والفيتوالكسينات. يحفز إنزيم ال Anthranilate synthase التفاعل الأول في المسار المتعدد الخطوات لتخليق التربتوفان عن طريق تحويل ال Chroismate إلى Anthranilate. يعد جهاز قاذف الجينات من إحدى الطرق التي تستخدم لإنتاج ذرة معدل وراثيا حيث تم استخدام البلازميد-pC2ASA2-NOS

ASA2&β)ASB with (feedback anthranilate synthase)ولذي يحمل جينات المسنونة عن زيادة الحمض الأميني التريبتوفان تحت التحكم من البادئCaMV35S. وضع الكالس علي بيئة ذات إسموزية عالية لمدة أربعة ساعات قبل قذف الكالس بالبلازميدات المحمّنة علي حبيبات الذهب. تم إستخدام نوعان من ضغط الهيليوم 900 رطل/ بوصة², 1100 رطل/ بوصة². وتم القذف إما مرة أو مرتان متتاليتان. ثم وضع الكالس علي بيئة إختيارية تحتوي علي 6MT 100 ميكرومول من المشبط. تم إستخدام ال Southern blotting , PCR لتحديد الجينات ومعرفة النباتات المحورة وراثيا وتم الحصول علي نباتات محورة وراثيا والتي أظهرت زيادة في الحمض الأميني التريبتوفان من 12-16 ضعف مقارنة بالنباتات غير المحورة وراثيا. كذلك ووجد أن كفاءة قذف الكالس مرتان أعلي من قنفة مرة واحدة بالبلازميد في إنتاج نباتات محورة وراثيا .