



Development of Multigenic Sugarcane to Decrease the Production Cost in Pakistan

Zahida Qamar^{1*}, Idrees Ahmad Nasir¹, Mounir G Abouhaidar², Kathleen L Hefferon³ and Ahmad Ali Shahid⁴

¹Centre of Excellence in Molecular Biology, University of the Punjab Lahore, Pakistan

²Department of Cell and Systems Biology, University of Toronto, Canada

³Department of Microbiology, Cornell University Ithaca, USA

⁴Institute of Molecular Biology and Biotechnology, The University of Lahore, Pakistan

Abstract

Due to higher amount of sucrose, sugarcane is grown commercially. In order to save sucrose yields, various studies have been designed to develop resistance in sugarcane against weeds and stem borers. In this study, two problems had been addressed by genetic manipulation of sugarcane to make them resistant against both herbicides and insects by expressing glyphosate resistant gene (CEMB-GTGene) and borer resistant genes (CEMB-Cry1Ac and CEMB-Cry2A) under control of NOs terminator and maize ubiquitin promoter. Mortality percentage of shoot borers *Chilo infuscatellus* was determined by assessing the Cry proteins through insect Bio-toxicity assays. Results showed that in 80 days old transgenic plants, 100% mortality rates of *Chilo infuscatellus* have been found showing that there was high resistance in transgenic sugarcanes against shoot borers and sufficient gene expression to fully resist target pests. Weed management was done by glyphosate spray assays. 70% to 76% of the transgenic plants were identified to be glyphosate resistant (3000 ml/Ha) in V₁ generation while 100% tolerant in V₂ generation. Thus, this transgenic sugarcane will help to boost sugarcane yield in the country as it now successfully provides resistance against both stemborers and glyphosate herbicides.

Introduction

Sugarcane is considered to be the world's significant cash crop as it is being cultivated around the globe in 58 countries [2] and 26.9 million hectares of area is used for sugarcane cultivation worldwide [1]. The 80% of world's sugar need is fulfilled by sugarcane *via* chemically synthesized sweetener known as sucrose [3]. A wide range of products are obtained from sugarcane like chemicals, biofuels, fibers, paper, beverages, detergents, insecticides, industrial enzymes, plastics, paints, pharmaceutical products, synthetics, chipboard and industrial chemicals like dextran, furfural and alcohol [4]. Sugarcane contributes to 0.7% GDP and 3.4% of agriculture sector and is cultivated on ~1.3 million hectare area in Pakistan [5]. 37% of the agriculture production in Pakistan is lost out of which 13% is because of insects [6]. Sugarcane crop is destroyed by ~1300 insect pests all over the world and by 61 insect species in Pakistan [7]. In Pakistan, 15% to 36% of sugarcane yield is lost due to stemborers, 10% to 20% by root-borers and 10% to 15% by top-borers [8]. Main objective of this study to prevent yield loss by making sugarcane resistant against stemborers and herbicides.

Materials and Methods

A gene cassette was designed that contains herbicides and stemborers resistant genes i.e. CEMB-GTGene, CEMB-Cry1Ac and CEMB-Cry2A under control of NOs terminator and maize ubiquitin promoter. These constructs pCEMB-SGTG and pCEMB-SC12 were introduced *via* electroporation into the *Agrobacterium* cells [9]. Colony PCR was performed to confirmed gene transformation *via* gene specific primers. 8 to 10 weeks old leaves of tobacco plants were co-cultured with *Agrobacterium* to induce *Agrobacterium* mediated transformation [10]. Expression of transgenes was indicated by histochemical detection of the GUS activity that was used as a reporter using agroinfiltrated leaves. Biolistic transformation method was used for transformation of transgene in 4 sugarcane varieties i.e. CPF-246, HSF-240, SPF-234 and SPF-213 [11]. During early transgenesis, transgene expression was determined by performing GUS assays on young shoots. Presence of transgenes was

OPEN ACCESS

*Correspondence:

Zahida Qamar, Centre of Excellence in Molecular Biology, University of the Punjab Lahore, Pakistan,
E-mail: Zahida.Qamar@cemb.edu.pk

Received Date: 30 Nov 2021

Accepted Date: 30 Dec 2021

Published Date: 04 Jan 2022

Citation:

Qamar Z, Nasir IA, Abouhaidar MG, Hefferon KL, Shahid AA. Development of Multigenic Sugarcane to Decrease the Production Cost in Pakistan. *Am J Leuk Res.* 2022; 5(1): 1022.

Copyright © 2022 Zahida Qamar. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

confirmed through PCR screening using CEMB-GT*Gene*, CEMB-Cry1Ac and CEMB-Cry2A genes specific primers. Stable transgene integration was determined by performing southern blotting [12] on PCR positive transformants. CEMB-GT*Gene*, CEMB-Cry1Ac and CEMB-Cry2A genes expression were determined through dipstick assays that were coated with the IgG monoclonal antibodies for each gene. These sticks were dipped in total proteins that were isolated from the fresh transgenic plants leaves [13]. ELISA was performed to quantify the transgene expressions. Toxicity effects of CEMB-Cry1Ac and CEMB-Cry2A endotoxins were determined by performing leaf biotoxicity assay on the leaves. CEMB-GT*Gene* expression and activity was confirmed by spraying glyphosate on the transgenic plants. Comparison between different lines (control and transgenic) was done through statistical analysis (Dunnett's tests, LSD and ANOVA).

Results

Restriction of the CEMB-GT*Gene*, CEMB-Cry1Ac and CEMB-Cry2A genes generated 1.4 kb, 1.8 kb and 1.9 kb fragments respectively which were then integrated into expression vector (pCAMBIA-1301). These constructs were introduced *via* electroporation into *Agrobacterium*. PCR analysis confirmed presence of transgenes. PCR positive transformants were subjected to agro-filtration using tobacco leaves in presence of GUS receptor. The expression of GUS was confirmed by bluish green color under fluorescent microscope. For sugarcane transformation and tissue culturing, CPF-246, HSF-240, SPF-234 and SPF-213 varieties of sugarcane were selected. To obtain

maximum embryogenic calli from the selected varieties, 4 different combinations were used for callus induction media. Maximum embryogenic calli was observed in CPF-246 (100%) followed by SPF-213 (90%), SPF-234 (90%) and HSF-240 (81%). Plasmid constructs were then transferred to these varieties *via* biolistic methods. Total of 400 explants were used for transformation. On single selection media (kanamycin), 91% of CPF-246, 74% of SPF-234, 70% of SPF-213 and 45% of HSF-240 survived while on double selection media (glyphosate and kanamycin), 81% of CPF-246, 40% of SPF-234, 34% of SPF-213 and 29% of HSF-240 transformed calli survived. Then after it, GUS assay was performed to screen the transgenic putative plants. PCR, southern blotting, dipstick assay and ELISA was performed for transgenic plants at V₀, V₁ and V₂ generation. Leaf bioassay was performed to determine the efficiency of CEMB-GT*Gene*, CEMB-Cry1Ac and CEMB-Cry2A genes and 60% to 100% mortality rate *Chilo infuscatellus* was determined in transgenic leaves. Weed management was done by glyphosate spray assays and 70% to 76% of the transgenic plants were identified to be glyphosate resistant (3000 ml/Ha) in V₁ generation while 100% tolerant in V₂ generation (Figure 1).

Discussion

Main objective of crop production is to obtain high yields even for sugarcane [14]. Different viruses, drought stress, weeds and insects are the major constrains for sugarcane [15]. Present study aimed to control insects and weeds through genetic manipulation of sugarcanes. In this study, for maximum callus regeneration, an efficient procedure

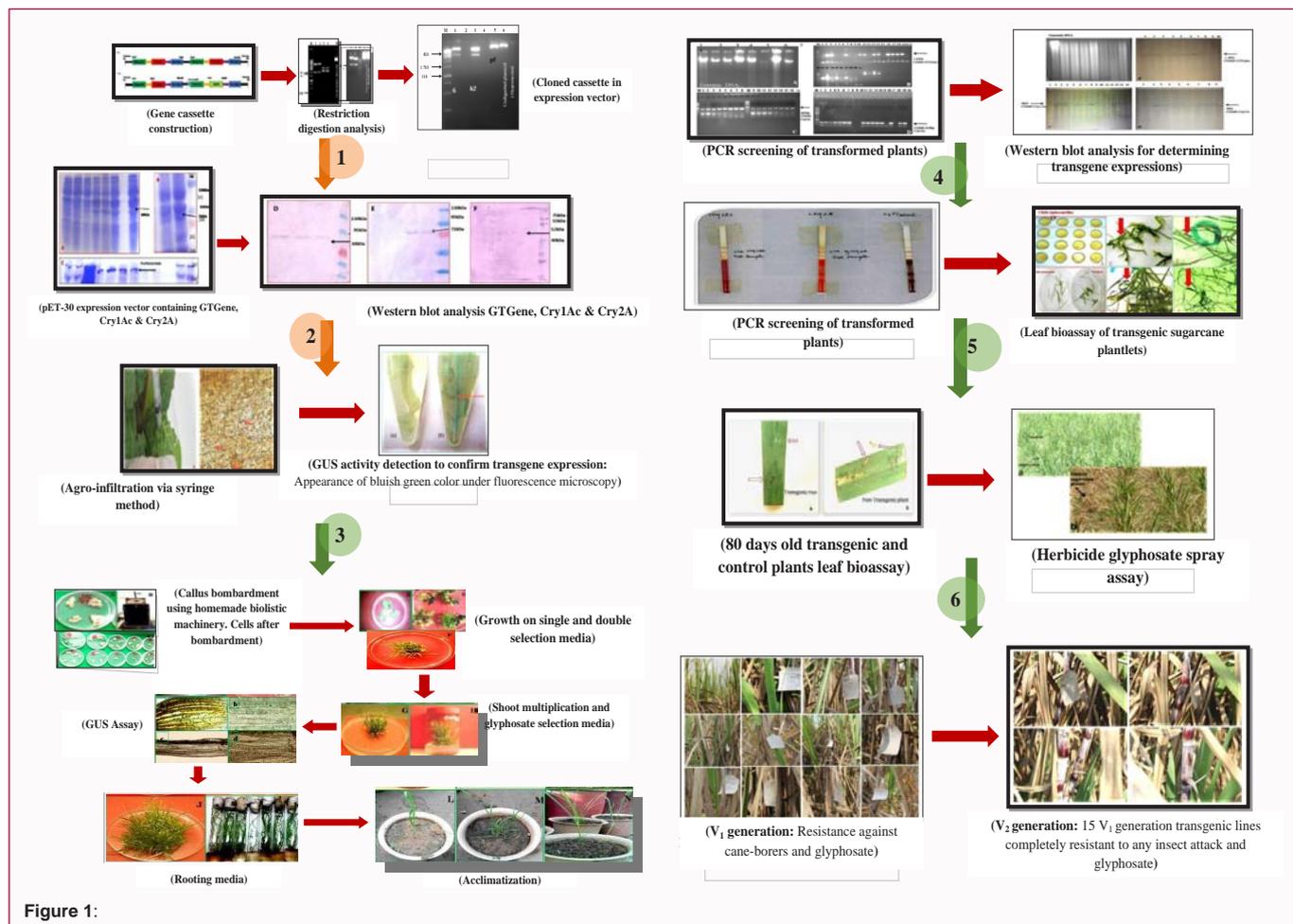


Figure 1:

was developed to instill tolerance against glyphosate and cane borers. For embryogenic callus formation, immature leaves were found to be excellent explants [16]. It basically strengthens the procedure for gene transformation in sugarcane [17]. From a callus inducing media with 2,4-D, embryogenic calli were obtained for all 4 varieties [18]. To enhance potential of embryogenic calli of sugarcane, it was supplemented with casein [19]. Tissue culture response was observed to critically screen all 4 varieties [20]. For genetic modification, varieties were selected on basis of regeneration response [21]. Studies have also disclosed that resistant against lepidopteran insects were best provided by Cry proteins [22]. Most commonly used herbicide for weed control is glyphosate which is a broad spectrum herbicide. One of the main drawbacks of using glyphosate is that along with weeds and herbs, it also stunts the plant growth thus affecting its yield [23]. It inhibits formation of EPSPS enzyme in shikimate pathway which leads to shikimate pathway being shut down. This inhibits the formation of 3 essential amino acids i.e. phenylalanine, tyrosine and tryptophan that humans can't synthesize and is required from plant source [24]. In this study, resistant against glyphosate and stem borers are provided by introducing CEMB-*GT*Gene, CEMB-*Cry1Ac* and CEMB-*Cry2A* genes into the sugarcane varieties.

Conclusion

In this study, 100% mortality rates of *Chilo infuscatellus* have been found in CPF-246 variety showing that there was high resistant in transgenic sugarcane against shoot borers and sufficient gene expression to fully resist target pests. Weed management was done by glyphosate spray assays and 70% to 76% of the transgenic plants were identified to be glyphosate resistant (3000 ml/Ha) in V₁-generation while 100% tolerant in V₂-generation. This study reported that after approval from biosafety committee, farmers can use this sugarcane variety as starting material for cost effective weeds and insect's control. More studies should be done to enhance stable *Bt*. toxin expression. Glyphosate resistant crops against 5000 mL/ha were recommended to be successful in controlling all sugarcane weeds.

References

1. Yao W, Ruan M, Qin L, Yang C, Chen R, Chen B, et al. Field performance of transgenic sugarcane lines resistant to sugarcane mosaic virus. *Front Plant Sci.* 2017;8:104.
2. Mayavan S, Subramanyam K, Jaganath B, Sathish D, Manickavasagam M, Ganapathi A. Agrobacterium-mediated in planta genetic transformation of sugarcane setts. *Plant Cell Rep.* 2015;34(10):1835-48.
3. Gao S, Yang Y, Wang C, Guo J, Zhou D, Wu Q, et al. Transgenic sugarcane with a cry1Ac gene exhibited better phenotypic traits and enhanced resistance against sugarcane borer. *PLoS One.* 2016;11(4):e0153929.
4. Raghavi S, Sindhu R, Binod P, Gnansounou E, Pandey A. Development of a novel sequential pretreatment strategy for the production of bioethanol from sugarcane trash. *Bioresour Technol.* 2016;199:202-10.
5. Farooq O. Agriculture, economic survey of Pakistan. *PBoS.* 2015.
6. Awan MF, Ali A, Muzaffar A, Abbas MA, Rao AQ, Qamar Z, et al. Transgenic cotton: Harboring board term resistance against insect and weeds through Incorporation of CEMB double Bt and cp4EPSPS genes. *Pak J Agri Sci.* 2016;53(3):501-5.
7. Long WH, Hensley SD. Insect pests of sugar cane. *Annu Rev Entomol.* 1972;17(1):149-76.
8. Bhatti I, Panhwar D, Unar G, Chohan M, Gujar N, Panhwar M, et al. Incidence and intensity of borer complex infestation on different sugarcane genotypes under agro-climatic conditions of Thatta [2008]. *Pak J Sci.* 2009;60(3-4):103-06.
9. Qamar Z, Aaliya K, Nasir IA, Farooq AM, Tabassum B, Qurban A, et al. An overview of genetic transformation of glyphosate resistant gene in *Zea mays*. *Nat Sci.* 2015;13(3):80-90.
10. Bhaskar PB, Venkateshwaran M, Wu L, Ané JM, Jiang J. Agrobacterium-mediated transient gene expression and silencing: A rapid tool for functional gene assay in potato. *PLoS One.* 2009;4(6):e5812.
11. Nasir IA, Tabassum B, Qamar Z, Javed MA, Tariq M, Farooq AM, et al. Herbicide-tolerant sugarcane (*Saccharum officinarum* L.) plants: An unconventional method of weed removal. *Turk J Biol.* 2014;38(4):439-49.
12. Edwards K, Johnstone C, Thompson C. A simple and rapid method for the preparation of plant genomic DNA for PCR analysis. *Nucleic Acids Res.* 1991;19(6):1349.
13. Qamar Z, Riaz S, Nasir IA, Ali Q, Husnain T. Transformation and evaluation of different transgenic lines for Glyphosate tolerance and cane borer resistance genes in sugarcane (*Saccharum officinarum* L.). *Cytol Genet.* 2017;51(5):401-12.
14. Ali A, Muzaffar A, Awan MF, Din S, Nasir IA, Husnain T. Genetically modified foods: engineered tomato with extra advantages. *Adv Life Sci.* 2014;1(3):139-52.
15. Thiebaut F, Grativol C, Carnavale-Bottino M, Rojas CA, Tanurdzic M, Farinelli L, et al. Computational identification and analysis of novel sugarcane microRNAs. *BMC Genomics.* 2012;13(1):1-14.
16. Snyman S, Meyer G, Richards J, Haricharan N, Ramgareeb S, Huckett BI. Refining the application of direct embryogenesis in sugarcane: Effect of the developmental phase of leaf disc explants and the timing of DNA transfer on transformation efficiency. *Plant Cell Rep.* 2006;25(10):1016-23.
17. Fitch M, Lehrer AT, Komor E, Moore PH. Elimination of sugarcane yellow leaf virus from infected sugarcane plants by meristem tip culture visualized by tissue blot immunoassay. *Plant Pathol.* 2001;50(6):676-80.
18. Nawaz M, Ullah I, Iqbal N, IQBAL MZ, Javed MA. Improving *in vitro* leaf disk regeneration system of sugarcane (*Saccharum officinarum* L.) with concurrent shoot/root induction from somatic embryos. *Turk J Biol.* 2013;37:726-32.
19. Joyce P, Hermann S, O'Connell A, Dinh Q, Shumbe L, Lakshmanan P. Field performance of transgenic sugarcane produced using Agrobacterium and biolistics methods. *Plant Biotechnol J.* 2014;12(4):411-24.
20. Ali K, Raza G, Mukhtar Z, Mansoor S, Asad S. Ideal *in-vitro* culture and selection conditions for sugarcane genetic transformation. *Pak J Agri Sci.* 2015;52(1):43-9.
21. Bakhsh A, Rao AQ, Shahid AA, Husnain T. Spatio temporal expression pattern of an insecticidal gene (*cry2A*) in transgenic cotton lines. *Not Sci Biol.* 2012;4(4):115-9.
22. Riaz N, Husnain T, Fatima T, Makhdoom R, Bashir K, Masson L, et al. Development of Indica Basmati rice harboring two insecticidal genes for sustainable resistance against lepidopteran insects. *S Afr J Bot.* 2006;72(2):217-23.
23. Nawaz A, Haseeb A, Malik H, Ali Q, Malik A.. Genetic association among morphological traits of *Zea mays* seedlings under salt stress. *Biol Clin Sci Res J.* 2020;2020(1).
24. Castle LA, Siehl DL, Gorton R, Patten PA, Chen YH, Bertain S, et al. Discovery and directed evolution of a glyphosate tolerance gene. *Science.* 2004;304(5674):1151-4.