

## 612/E2

## Development of transgenic rice plants with lysine rich protein coding gene

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Even though it is the staple food of more than half the world's population, rice is not considered a rich source of protein. One of the essential amino acids, lysine is low in rice seed. Hence, this study was aimed at increasing both the lysine and total protein content in rice seeds by introducing the pollen-specific lysine-rich protein encoding gene (SBgLR) from potato (Solanum tuberosum) into indica rice (Oryza sativa L.) seed under the control of the rice seed-specific globulin promoter. Total Ribo Nucleic Acid (RNA) from potato pollen grains was extracted and the produced CDNA and SBgLR gene was amplified by Polymerase Chain Reaction (PCR) using SBgLR gene-specific primers. The isolated rice genomic Deoxy Ribonucleic Acid (DNA) from rice variety Bg 94-1 was subjected to PCR to amplify the promoter sequence of the globulin gene. The amplified promoter region was cloned into pGEM®-T Easy vector, and then into pCAMBIA1391Z vector. Recombinants were selected and sequenced. The SBqLR gene containing recombinant vector (pCR<sup>®</sup>2.1-TOPO-SBgLR), previously cloned in our laboratory was restriction digested with BamH1 and EcoR1 enzymes, and cloned into the corresponding sites of pCAMBIA1391Z-Glb vector construct. Electro competent cells of Agrobacterium strain GV3101 was transformed with the recombinant construct (pCAMBIA1391Z-Glb-SBqLR). Bq 94-1 rice seed derived calli, and active rice embryos were used for transformation. The embryo transformation method proved less time consuming and more effective in producing transgenic rice plants. PCR analysis of regenerated transformed plants indicated the presence of the SBgLR gene.

Keywords: Agrobacterium, lysine, transgenic rice, transformation

Acknowledgement: TG11/53 from the Sri Lanka Treasury