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Introgression of Bacterial Blight Resistance Genes (*Xa*21, *xa*13 and *xa*5) into CB 174 R, an Elite Restorer Line in Rice

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Abstract: Bacterial blight caused by *Xanthomonas oryzae* pv. *oryzae* (Xoo) is one of the major disease caused severe yield reduction in the rice growing regions. One dominant (*Xa*21) and two recessive genes (*xa*13 and *xa*5) were introgressed into CB 174 R through marker assisted breeding. The present study found three (*Xa*21+*xa*13+*xa*5) and two (*Xa*21+*xa*13 or *Xa*21+*xa*5 or *xa*5+*xa*13) genes introgressed combinations in the early segregated materials through foreground selection. The identified homozygous/heterozygous individuals forwarded to next cycles of breeding to fix homozygous conditions for all three genes with an improved agronomic performance background, and thus could be used as a donor source for future rice breeding programme.

Keywords: marker assisted selection; gene specific marker; gene pyramiding

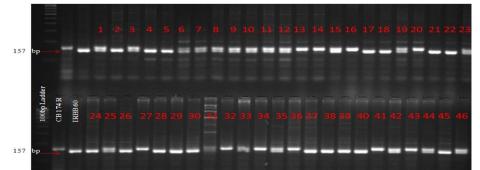
Results and Discussion

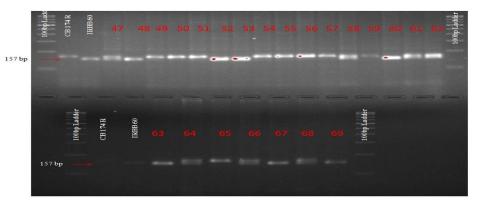
- ➢ Fifty-four (42 having *Rf*4) out of 110 F₂ individuals identified as having three/two genes combination in this study.
- Five F_2 individuals were in heterozygous state for all three genes (*Xa5xa5*, *Xa13xa13* and *Xa21Xa21*), and also two individuals had heterozygous resistance for two loci (*Xa5xa5* and *Xa13xa13*) and homozygous for one locus (*Xa21Xa21*), along with fertility genes which were characterized earlier.
- Presence of Xa21 in homozygote or heterozygote state in combinations with other genes found to have higher level of resistance.
- Also, two recessive genes shown higher level of resistance when they were in homozygote (*xa5xa5* and *xa13xa13*) than heterozygote (*Xa5xa5* and *Xa13xa13*) condition.
- ▶ The identified different combinations of homozygous/heterozygous resistance plants F_2 with fertility restoration genes, and the subsequent $F_{2:3}$ families showed an improved agronomic performance would be used as a donor parent for future rice breeding programme.

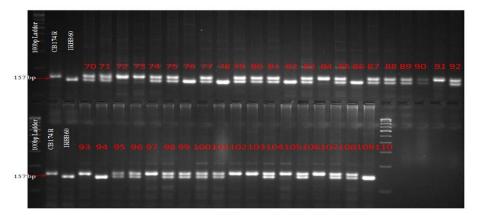
Conclusions

BB resistance genes identified in heterozygous and/ or homozgygous with superior agronomic performances of the studied breeding materials led to use as a donor parent in the BB resistance genes introgression breeding.

100bp Ladder CB 174 R IRBB 60







Identification of Xa21 gene in rice

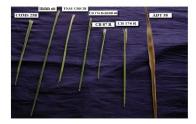


Bacterial inoculum



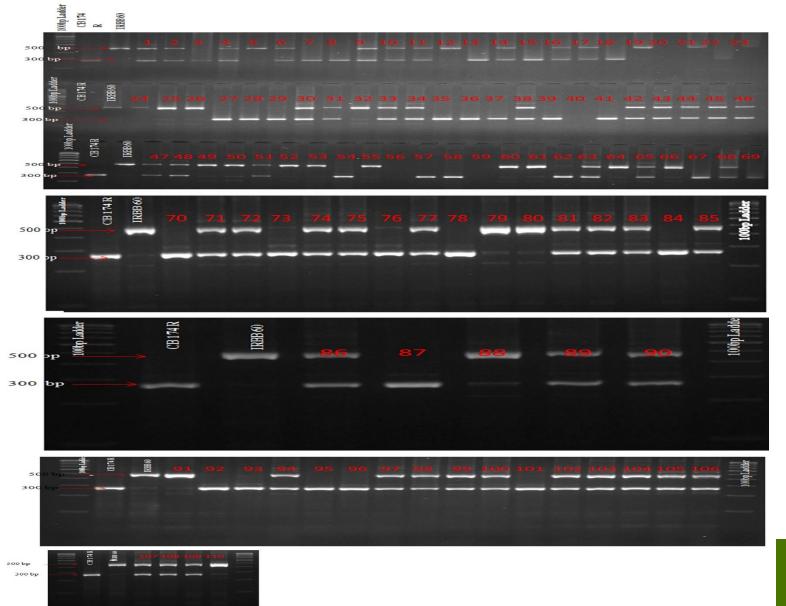
Inoculation with Spores by Leaf clipping method





Reaction of Parental lines against BB

Phenotype screening for BB in parents and F_1



Identification of *xa*13 gene in rice

Acknowledgments

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