

Advancement in molecular approaches for biotic stress tolerance in plants

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Introduction

'Stress' in plants can be defined as any external factor that negatively influences plant growth, productivity, reproductive capacity or survival. This includes a wide range of factors that can be broadly divided into two main categories: abiotic or environmental stress

factors and biotic or biological stress factors. **Biotic stress** occurs as a result of damage done to the plant by other living organisms such as bacteria, viruses, fungi, parasites, beneficial and harmful insects, weeds and cultivated or native plants (David *et al.* 2001).

Types of biotic stresses

Fungi Virus Bacteria and Nematode

Types of response in stress:

- 1) Reactive oxygen species
- 2) Calcium signaling
- 3) Hypersensitive response
- 4) Systemic acquired response
- 5) Salicylic acid

6) Jasmonic acid

7) PR Proteins

Strategies for imparting stress tolerance in plants: Conventional breeding, molecular breeding, genetic engineering and novel approaches

Review of research work

Sohrab *et al.* (2016) studied transgenic cotton plants that were developed by using β C1 gene in antisense orientation gene driven by Cauliflower mosaic virus-35S promoter and nos (nopaline synthase) terminator and mediated by *Agrobacterium tumefaciens* transformation and somatic embryogenesis system. The developed transgenic and inoculated plants remained symptomless till their growth period. the plants were observed as resistant to CLCuV.

Nekrasov *et al.* (2017) reported on Tomelo a non-transgenic tomato variety resistant to the powdery mildew fungal pathogen using the CRISPR/Cas9 technology. They used whole-genome sequencing to show that Tomelo does not carry any foreign DNA sequences but only carries a deletion that is indistinguishable from

naturally occurring mutations. They also presented evidence for CRISPR/Cas9 being a highly precise tool, as they did not detect off-target mutations in Tomelo.

Nevame *et al.* (2018) studied that TYLCV resistance has been based mostly on Ty-3 as a race-specific resistance gene by introgression originating from wild tomato species relatives. The newly developed marker was named ACY. The reliability and accuracy of ACY were evaluated against those of Ty-3 linked marker P6-25 through screening of commercial resistant and susceptible tomato hybrids, and genetic segregation using F2 population derived from a commercial resistant hybrid AG208. With the use of bioinformatics and DNA sequencing analysis tools, deletion of 10 nucleotides was observed

in Ty-3 gene sequence for susceptible tomato variety. ACY is a co-dominant indel-based marker that produced clear and strong polymorphic band patterns for resistant plants distinguishing it from its susceptible counterpart.

Liu *et al.* (2020) selected a panel of *Yr* gene pyramiding lines with Chuanyu12 as the background parent. The number of pyramided *Yr* genes was significantly correlated with

stripe rust resistance. *Yr15*, *Yr62*, and *Yr65* are effective to the current *Pst* races. Pyramiding more than four effective or partially effective *Yr* genes can provide enough resistance to stripe rust. Additive effects or epistatic effects existed in gene combinations in this study such as *Yr26 + Yr48*, *Yr30 + Yr64*, and *Yr30 + Yr48*. *Yr*-gene pyramiding lines with desirable agronomic traits were obtained for durable controlling *Pst* in wheat breeding.

Conclusions

- Conventional knowledge has almost saturated in finding the solution for the increasing biotic stress resulting due to climatic change and other causes.
- Genetic engineering has proved its worth in tweaking the plant's ability to cope with various stresses.
- CRISPR/Cas9 has the ability to generate specific double-stranded breaks and able to precise editing of the genome.
- The antisense approach has also been utilized, based on homology dependent for development of resistant transgenic plants against disease resistance.

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