## **Research Article: PR Verma Award for PhD Students – 2015 Runner**

## Fluorescent Pseudomonads/Abiotic Elicitor–Mediated Induced Systemic Resistance in Cabbage Against *Xanthomonas campestris* pv. *campestris* in Gangetic Alluvial Region of West Bengal

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## Abstract

Black rot of cabbage caused by Xanthomonas campestris py. campestris (Xcc) is a serious menace in the cultivation of cabbage and other cole crops in West Bengal. In the present investigation, attempts were made for determining the molecular variability of Xcc and induced systemic resistance mediated through fluorescent pseudomonads and abiotic elicitors were explored for the management of black rot disease of cabbage under Gangetic Alluvial Region of West Bengal. Thirty-one isolates of xanthomonads were isolated from cabbage and cauliflower and were confirmed through biochemical and molecular characterization using primers targeting hrpF homologous to nolX host recognition protein. All the thirty-one isolates were identified as Xcc. Sequencing of Internal Transcribed Spacer (ITS) regions identified the length of ITS 1, ITS 2 and ITS 3 as 112, 19 and 225 nucleotides. In silico phylogenetic study based on ITS sequence alignment could identify very high similarity among Xcc isolates and other xanthomonads such as X. campestris pv. raphani, X. arboricola pv. juglandis, X. cassava, X. pisi, X. oryzae pv. oryzae, X. vasicola and X. cucurbitae and differentiated seventeen number of Xanthomonas species into two major clusters. Studies on the ISR mediated through fluorescent pseudomonads isolates indicated that Pseudomonas aeruginosa isolates PT1 and PB3 collected from cabbage phyllosphere and rhizosphere were antagonistic to Xcc, Combined application of PT1 using seed treatment, rood dipping and foliar spray reduced the disease incidence by 50.6 per cent followed by foliar application of acibenzolar S-methyl (ASM) at 5 mM. It reduced the disease severity by 48.1 per cent and was not found to differ significantly from P. aeruginosa isolates. Changes in enzyme activities including phenol content were all negatively correlated with the disease progression. Under PT1 mediated induced systemic resistance (ISR) in cabbage-Xcc system, changes in peroxidase (POX) activity was significantly correlated with the disease progression. However, in ASM mediated resistance in cabbage-Xcc system, both phenylalanine ammonia lyase (PAL) and POX activity were significantly correlated with disease progression.

Key words: Biocontrol, isozymes, *Pseudomonas aeruginosa*, systemic acquired resistance, *Xanthomonas campestris* pv. *campestris* 

**Citation:** Barman Roy A and Dutta S. 2017. Fluorescent pseudomonads/abiotic elicitor-mediated induced systemic resistance in cabbage against *Xanthomonas campestris* pv. *campestris* in gangetic alluvial region of West Bengal. J Mycol Pl Pathol 47 (3): 262-274.