## Functional characterization of papain-like cysteine proteases genes in rice

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Papain-like cysteine proteases (PLCP) are key enzymes involved in cell death as response to biotic stress. Functional genetic investigation of cysteine protease family members has been performed in a fragmentary scale to understand its specific role in plants. Highlights of research milestone for these proteases provide strong evidence on their diverse and overlapping roles in basal immunity and effector-triggered immunity. The objective of this study was to provide useful insights into biological function of three cysteine protease genes, OsCP2, OsCP3, and OsCP5, in rice. Overexpression of rice cysteine protease attenuated the virulence of Xanthomonas oryzae pv. oryzae race K3a in all transgenic lines which displayed moderate resistance as indicated by shorter lesion lengths (OsCP2ox, 6.82 cm; OsCP3ox, 5.55 cm; and OsCP5ox, 5.40 cm) than wild type Dongjin (16.07 cm) whereas RNAi-mediated knockdown of OsCP3 resulted in severe bacterial leaf blight symptoms (17.1 cm). Abiotic screening revealed the biological significance of these three cysteine protease genes, especially of OsCP3, against salinity stress for which rice exhibited moderate tolerance (salinity score = 5.0 to 5.2). This study provides experimental evidence for roles of papain-like cysteine protease in improving resistance of rich against Xanthomonas oryzae pv. oryzae and tolerance against salinity stress, suggesting that these genes could be used as a valuable resource to be employed in rice breeding program to improve its ability to withstand biotic and abiotic stresses. Acknowledgements: This study was supported by the Russian Science Foundation. This work was supported by a grant from the Next-Generation BioGreen 21 Program (PJ01330201), RDA, Republic of Korea.