World Of Rudra Skyrim

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World of Rudra pc game, the chitinase gene was upregulated, and the activity of chitinase was detected in both plants (Maize and *C. comatus*) in response to the infection of CMD. This result indicated the important role of chitinase in the induction of defense responses and was consistent with the report of Yang et al. [@pone.0041518-Yang1]. In addition, the expression levels of genes regulated by wound-induced signal transduction network were also detected, such as genes related to MAPK, jasmonic acid (JA) and salicylic acid (SA). The results suggested that the increase of chitinase activity in *C. comatus* was accompanied with an increase of resistance to CMD. *C. comatus* did not elicit an elevation in ROS while maize

treated with the same pathogen rapidly increased the contents of ROS at the early stage of CMD infection. The present results indicated that ROS were detected in early stage of CMD infection while chitinase was activated in later stage of CMD infection. The accumulation of ROS and the activation of chitinase simultaneously may induce resistance to CMD in *C. comatus* while maize may respond to the resistance by activation of other signaling pathways such as SA pathway. Therefore, the activation of two pathways (MAPK and chitinase) could lead to the induction of resistance by *C. comatus* against CMD. In order to determine the gene regulation mechanism of *C. comatus* to CMD, a total of 220 CMD ESTs were used to perform the BLAST analysis of *C. comatus*. Three classes of transcription factors, including bHLH (basic

helix-loop-helix), GRF (growth-regulating factor) and WRKY (WRKY DNA-binding proteins), were obtained by BLAST search against the tomato EST database. Among these transcription factors, the bHLH class was found to be the most abundant and exhibited significant homology with well-characterized tomato bHLH proteins. In tomato, the activation of bHLH plays important roles in responses to the pathogens [@pone.0041518-Hussey1]. Moreover, we detected a bHLH (accession no. AC153094) that was isolated from *C. comatus* cDNA library, which was upregulated by CMD in *C. 6d1f23a050

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