Figure S1. Construction of Phylogenetic Tree by Combining Multiple Genes of HJGFB-1 strain. Phylogenetic tree inferred from the combined datasets (ITS, TEF, TUB, and RBP2) from members of the F. solani species analyzed. The evolutionary history was inferred by using the Bayesian Inference (BI) analyses, using *Fusarium plagianthi* NRRL22632 and *Fusarium vinguliforme* NRRL 32392 as the outgroup taxon.