



Figure S8: Phylogenetic tree of orthologues of QDE-1-like RNA-dependent RNA polymerase protein of *C. zeina* using representative fungal species from different orders in the class *Dothideomycetes*. *C. zeina* RdRp (PKS04443) is shown with a black dot. The tree was constructed using the Maximum Likelihood method in Mega-X, and was rooted using *Fusarium graminearum* and *Neurospora crassa* (class, *Sordariomycetes*) as an out-group. Bootstrap values greater than 50% are shown (1000 bootstraps). The scale bars indicate substitutions per site.