Figure S4: Phylogenetic tree of orthologues of the *C. zeina* DCL-1 protein using representative fungal species from different orders in the class *Dothideomycetes*. *C. zeina* DCL-1 (PKR99831) 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.