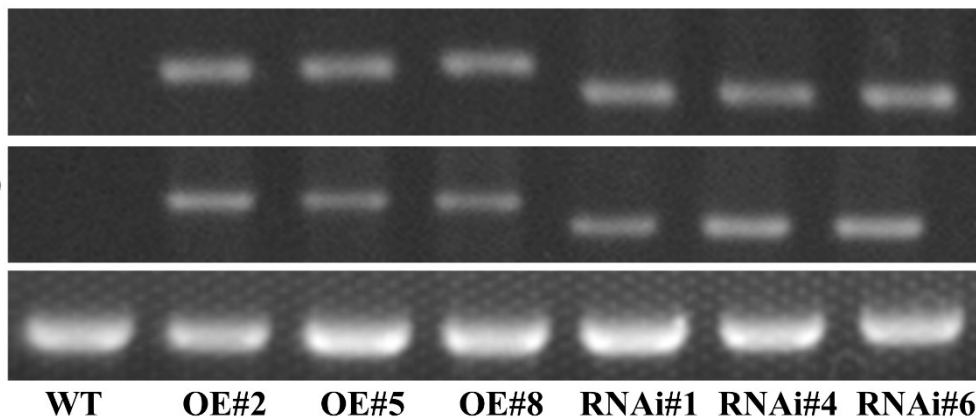**Figure 1.** Amino acid sequence alignment of BEC1019 homologs in different plant pathogens using DNAMAN software.

Figure 2. Semi-quantitative RT-PCR analysis of *BEC1019* expression in RNAi and overexpressing plants. *TaActin* transcripts were quantified as an internal control.