



Fig. S1. Bioinformatics analysis of *AsFADD*. (A) ORF sequence and amino acid sequence of FADD. The area marked with the start codon (ATG) and end codon (TGA) is the DED domain. The area underlined with the black font is the DED domain, and the area underlined with the red font is the Death domain. (B) Multiple comparisons of amino acid sequences of FADD and other species of FADD. (C) Phylogenetic tree of FADD amino acid sequence.