



Figure S1. The gene structure of *SmMFT-2* as predicted using Fgenesh gene prediction tool. CDSf indicates the first CDS (which starts with a start codon at 138th position in this image), CDSi indicates internal exon, CDSo indicates outer exon, and CDSl indicates the last coding sequence (which ends with a stop codon), TSS indicates Transcription Start Site and PolA indicates poly-A tail.