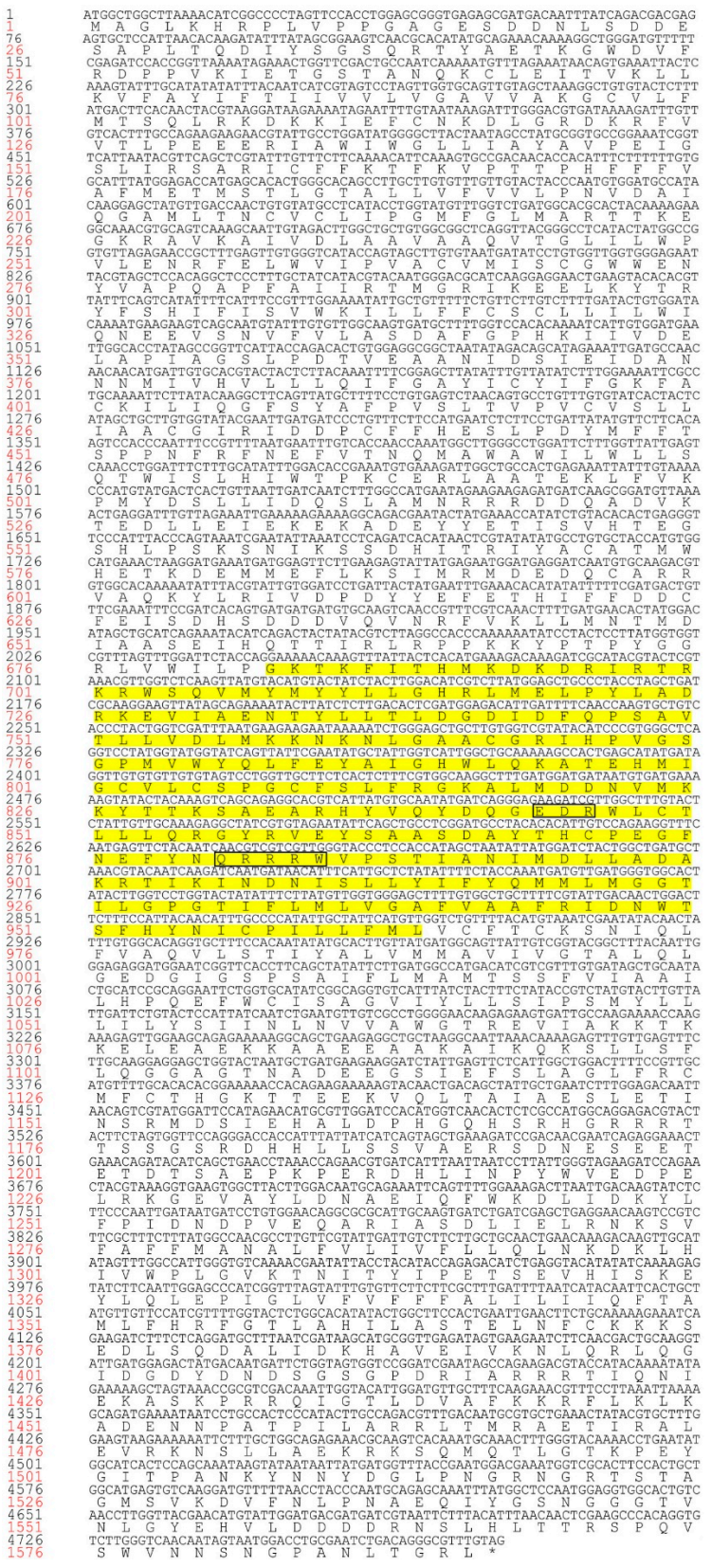

Figure S1 Nucleotide sequences for *L. trifolii* CHS2 cDNA and its predicted amino acid sequence. Nucleotide numbering starts with the adenine in the first methionine codon of the putative open reading frame. The highly conserved region, chitin_synth_2 domain, is highlighted. The asterisk indicates the translational termination codon. The EDR and QRRRW motifs are boxed.