Supplementary Figures

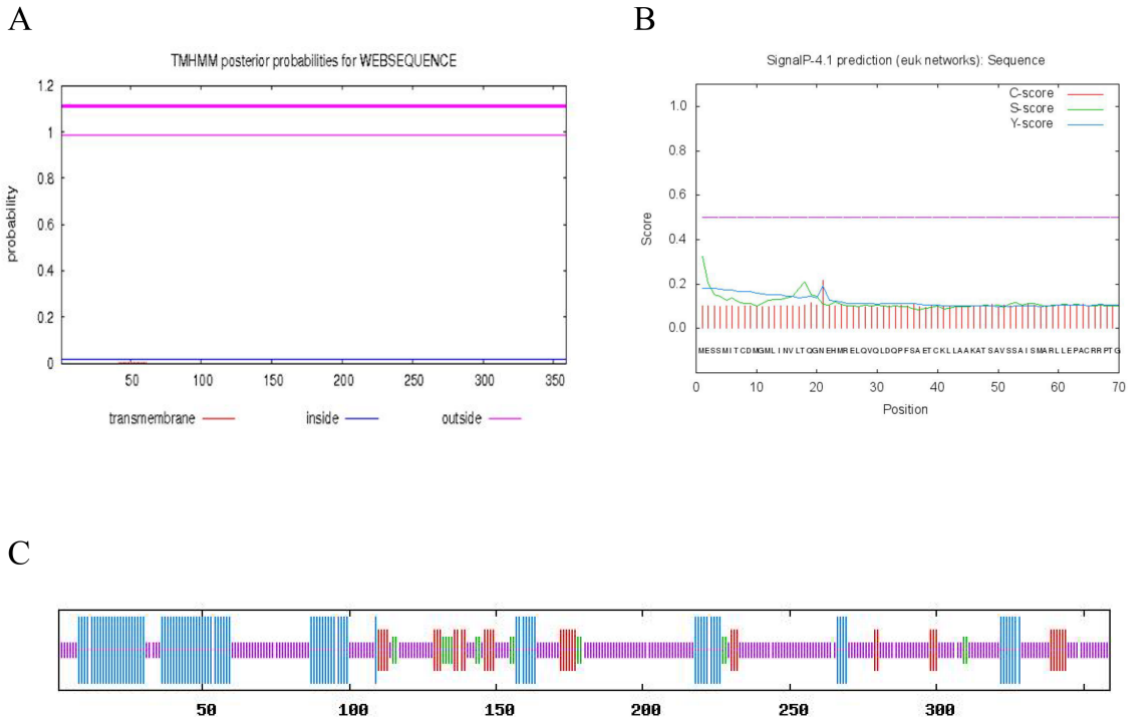

Figure S1: Bioinformatic analysis of CgWRKY53 protein structure.(A) Predicted transmembrane helical structure of CgWRKY53 protein; (B) Predicted signal peptide of CgWRKY53 protein;(C) Secondary structure prediction of CgWRKY53 protein: From left to right, representing the N-terminal to C-terminal regions. Blue indicates α-helix locations, red indicates extended chain locations, green indicates β-sheet locations, orange indicates random coil locations, and pink-purple indicates amino acids without specific structural features.