

Supplementary Materials

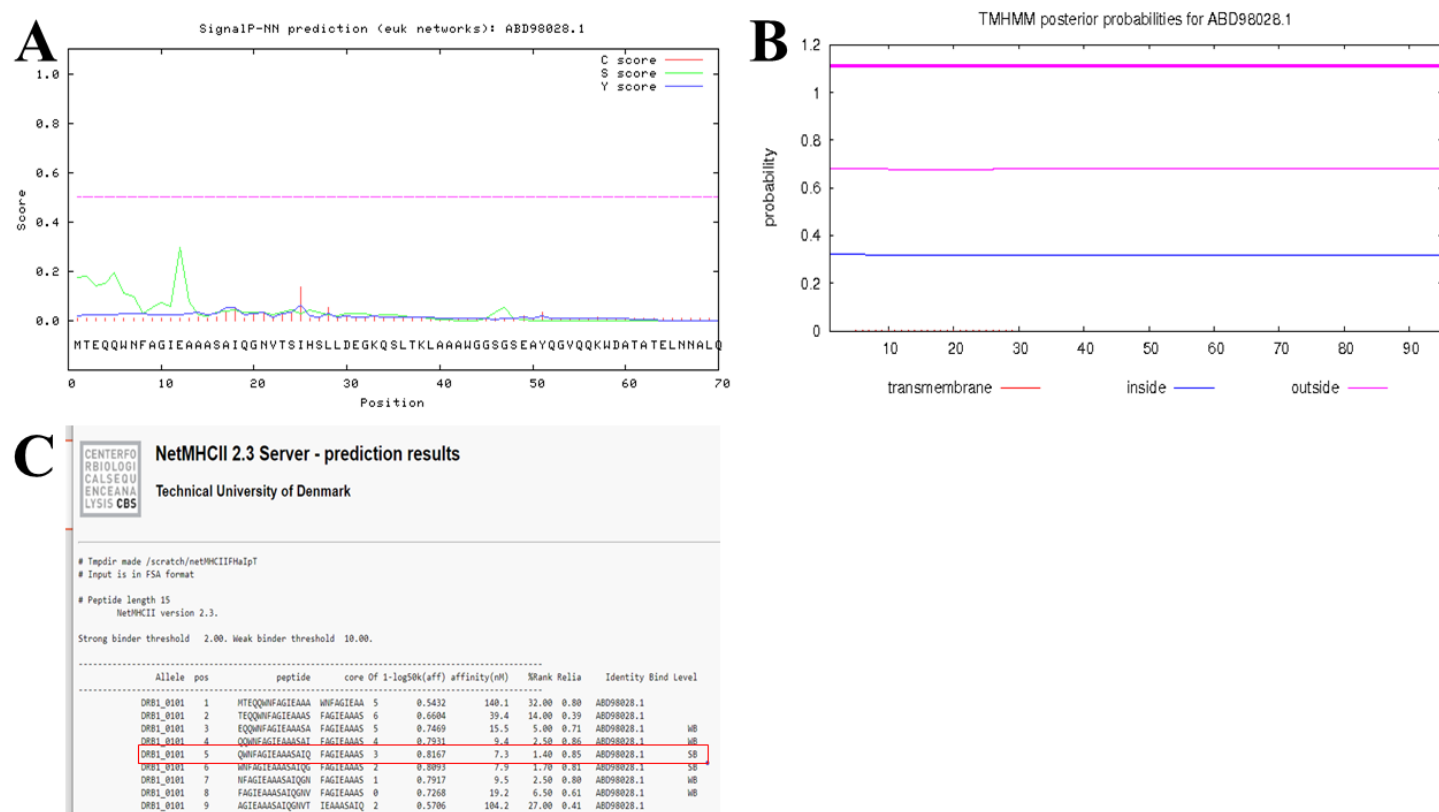


Figure S1. Analysis of the interaction of the ESAT-6 protein by integrating different algorithm searches. (A) Signal peptide prediction (SignalP peptide). (B). Prediction of transmembrane regions (TMHMM). (C). Prediction of epitopic regions for MHC II (NetMHCII).