Supporting Information:

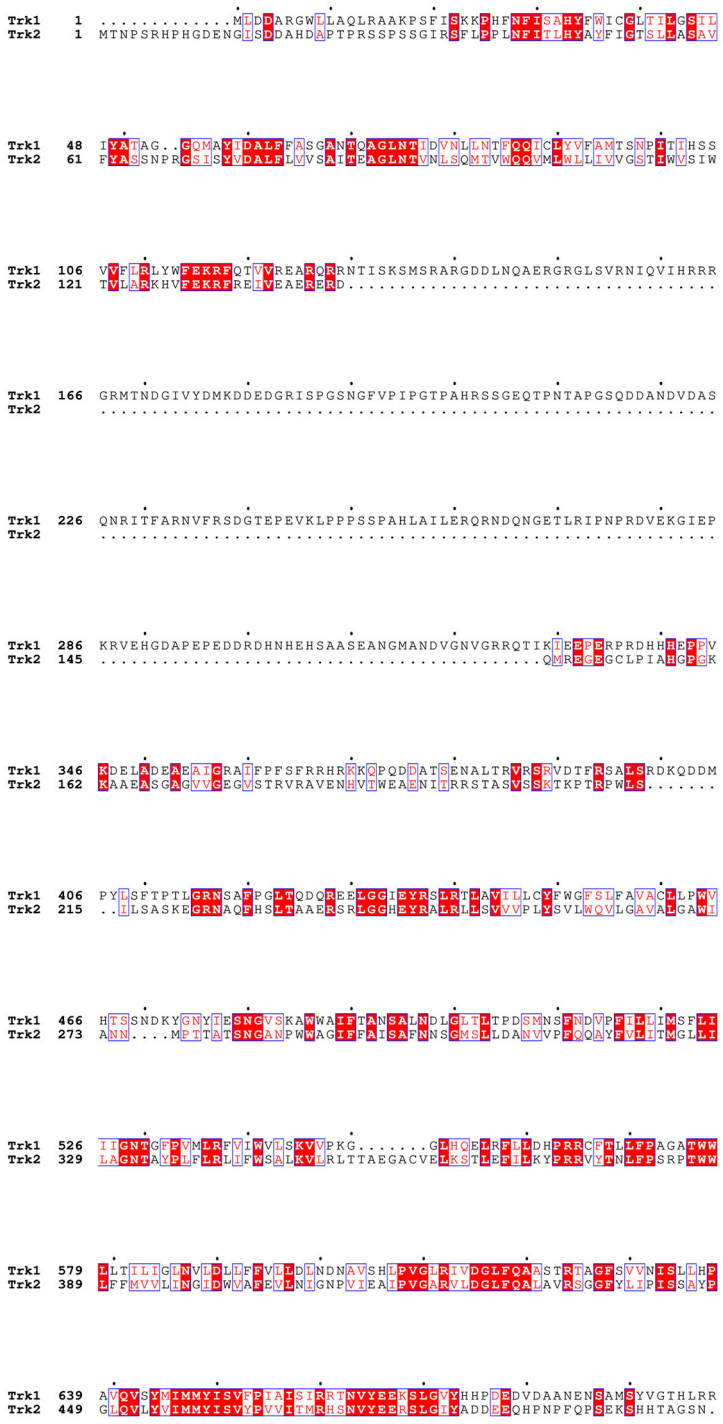

Figure S1 Amino-acid sequence alignment showed CgTrk1 and CgTrk2 were highly conserved.