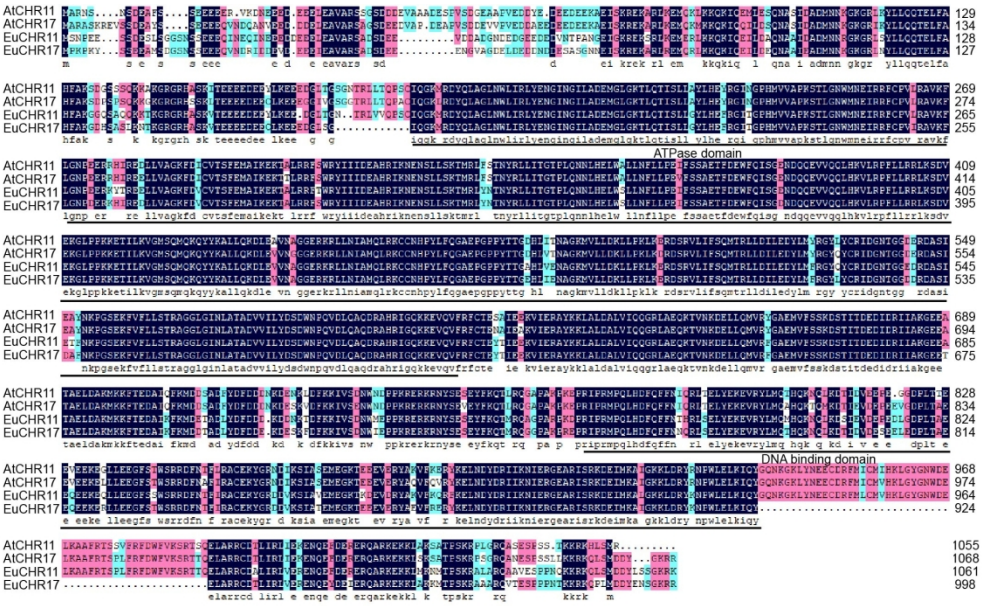

Supplemental Fig. S2 Amino acid sequences alignment of the CHR11 and CHR17 proteins in Arabidopsis and *E. ulmoides*. The conserved ATPase domain and DNA binding domain are labeled. Alignment was performed using DNAMAN software.