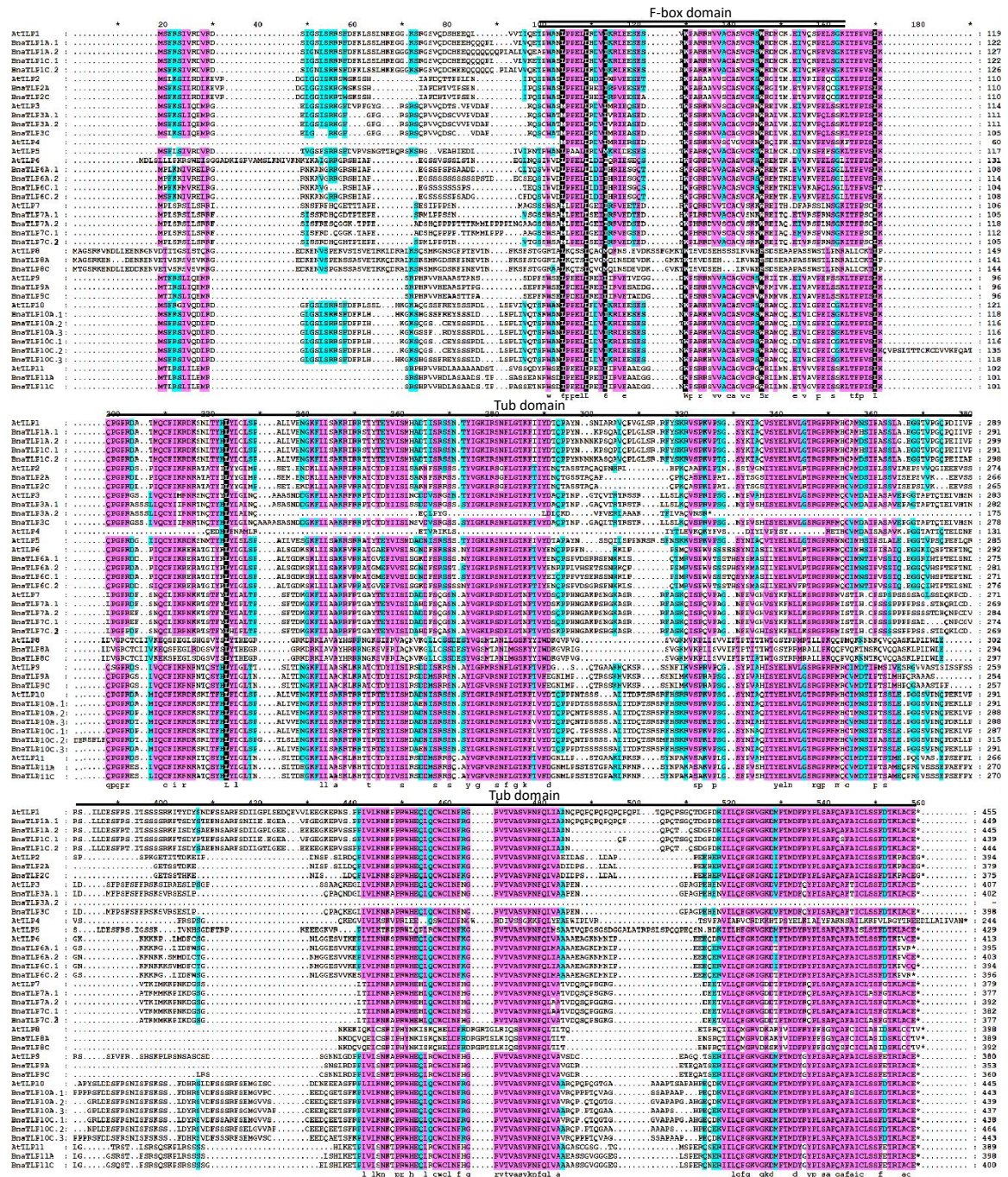




Supplementary Figure 1 | Multiple sequence alignment of BnaTLP proteins from *Brassica napus*. Black, pink and light blue shading, respectively, represent amino acids with 100, ≥ 75, and 50% similarity of amino acids. The locations of the F-box domain and Tub domain are indicated with double and single solid lines above the sequences, respectively. The alignment is generated by the ClustalW program.