

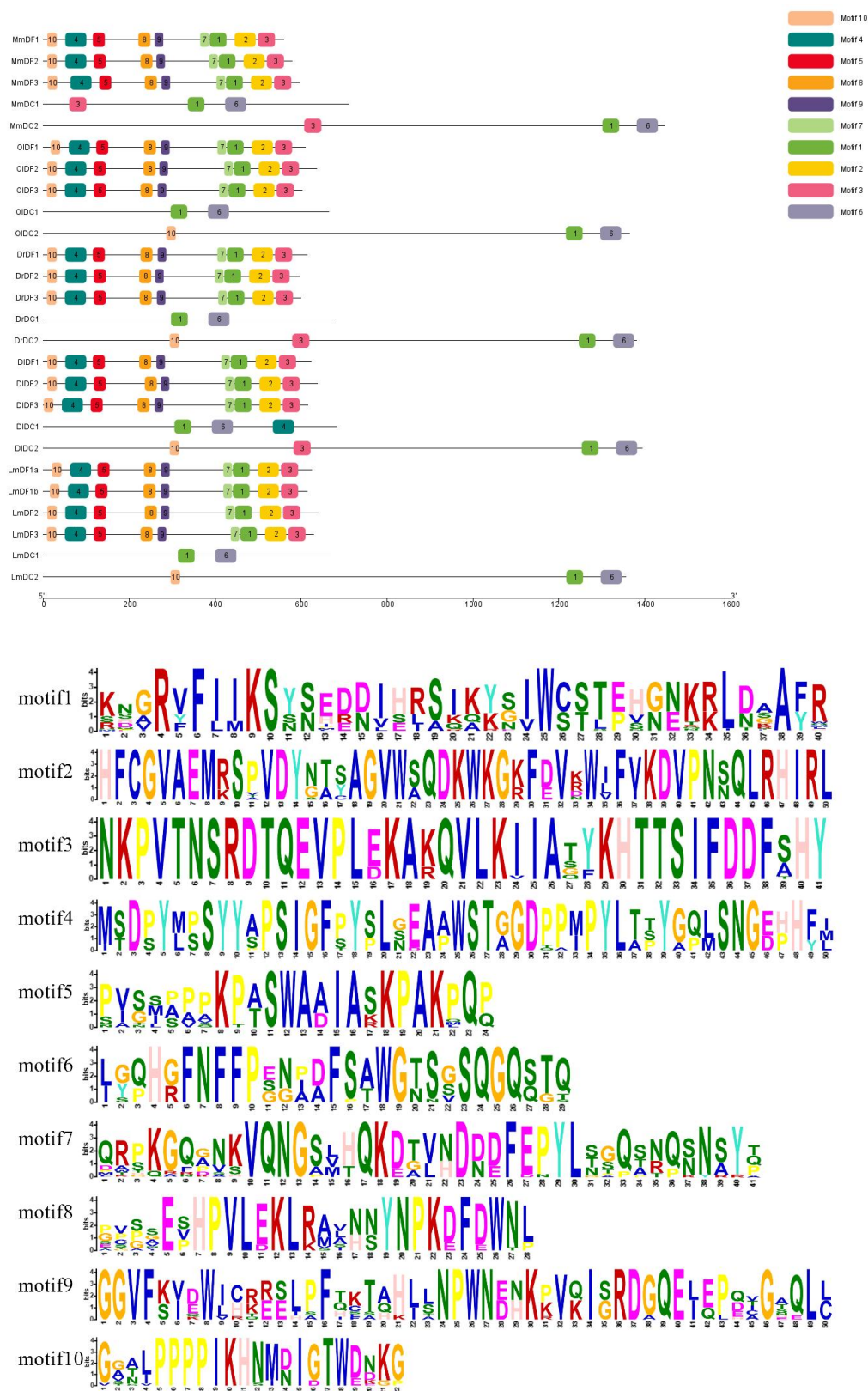


Fig. S2 Conserved motif analysis of *LmYTH* domain-containing genes.