



Fig. S2 Expression profiling of *GhMYB113*, *GhTT2-3A* and *GhbHLH130D* in different cotton tissues and developing fibers. (a) Expression of *GhMYB113* in different cotton tissues. Data are presented as mean \pm standard deviation (n=4). (b) Expression analysis of *GhMYB113*, *GhTT2-3A* and *GhbHLH130D* in fibers at 10, 15 and 18 DPA during development across different wild cotton accessions, including white cotton: Coker 312 (C312), J14 and brown cotton: Zong 2-63 (Z263), Bai991. The cotton ubiquitin gene *GhUBI1* was used as a reference gene for normalization. Data are presented as mean \pm standard deviation (n=4).