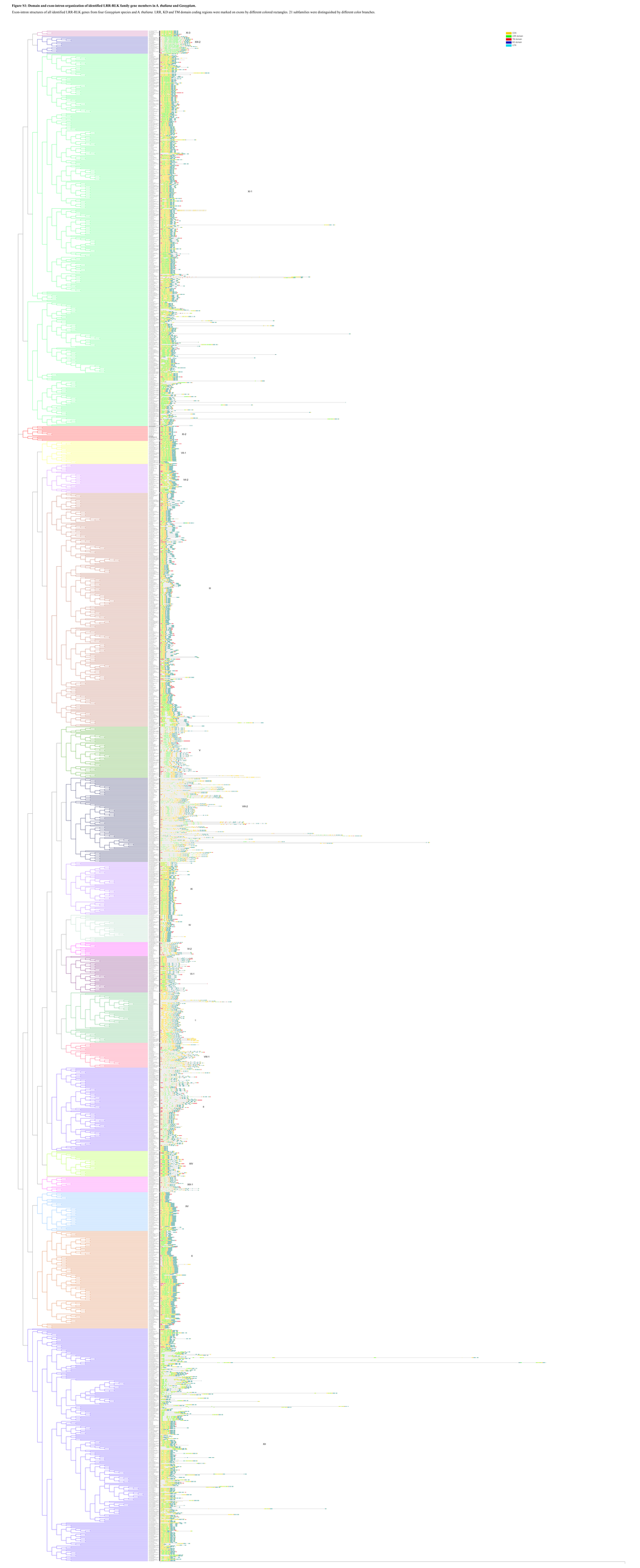
Figure S1: Domain and exon-intron organization of identified LRR-RLK family gene members in *A. thaliana* and *Gossypium*.

Exon-intron structures of all identified LRR-RLK genes from four *Gossypium* species and *A. thaliana*. LRR, KD and TM domain coding regions were marked on exons by different colored rectangles. 21 subfamilies were distinguished by different color branches.