

Full-length transcriptomic identification of R2R3-MYB family genes related to secondary cell wall development in *Cunninghamia lanceolata* (Chinese fir)

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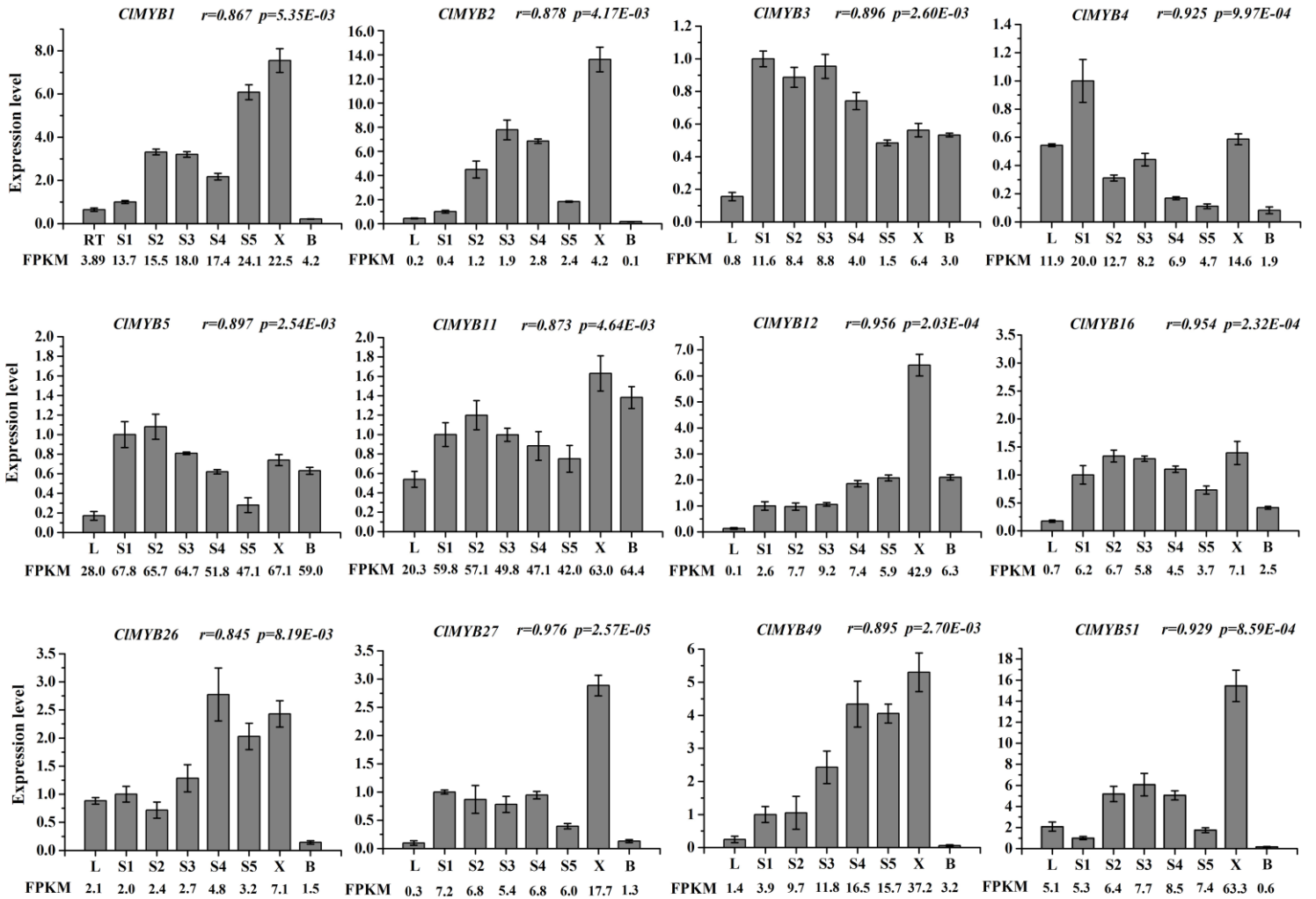


Figure S1. Expression profiles of 12 *CIMYB* genes in eight organs and tissues using qRT-PCR. Data were normalised to *ClActin* gene. The expression level in S1 was used as a reference and was set at 1. Error bars indicate standard deviation of three biological replicates. The corresponding FPKM values were listed. The r represented the correlation coefficient between qRT-PCR and RNA-Seq data, and the p was the P-value. L:leaves, S1-5: stem segments from top to bottom, X: xylem, B: bark