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

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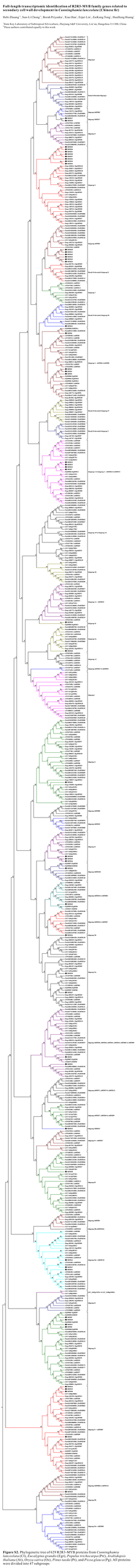

Figure S2. Phylogenetic tree of 629 R2R3-MYB proteins from *Cunninghamia lanceolata* (Cl), *Eucalyptus grandis* (Eg), *Populus trichocarpa* (Pt), *Arabidopsis thaliana* (At), *Oryza sativa* (Os), *Pinus taeda* (Pt), and *Picea glauca* (Pg). All proteins were divided into 47 subgroups.