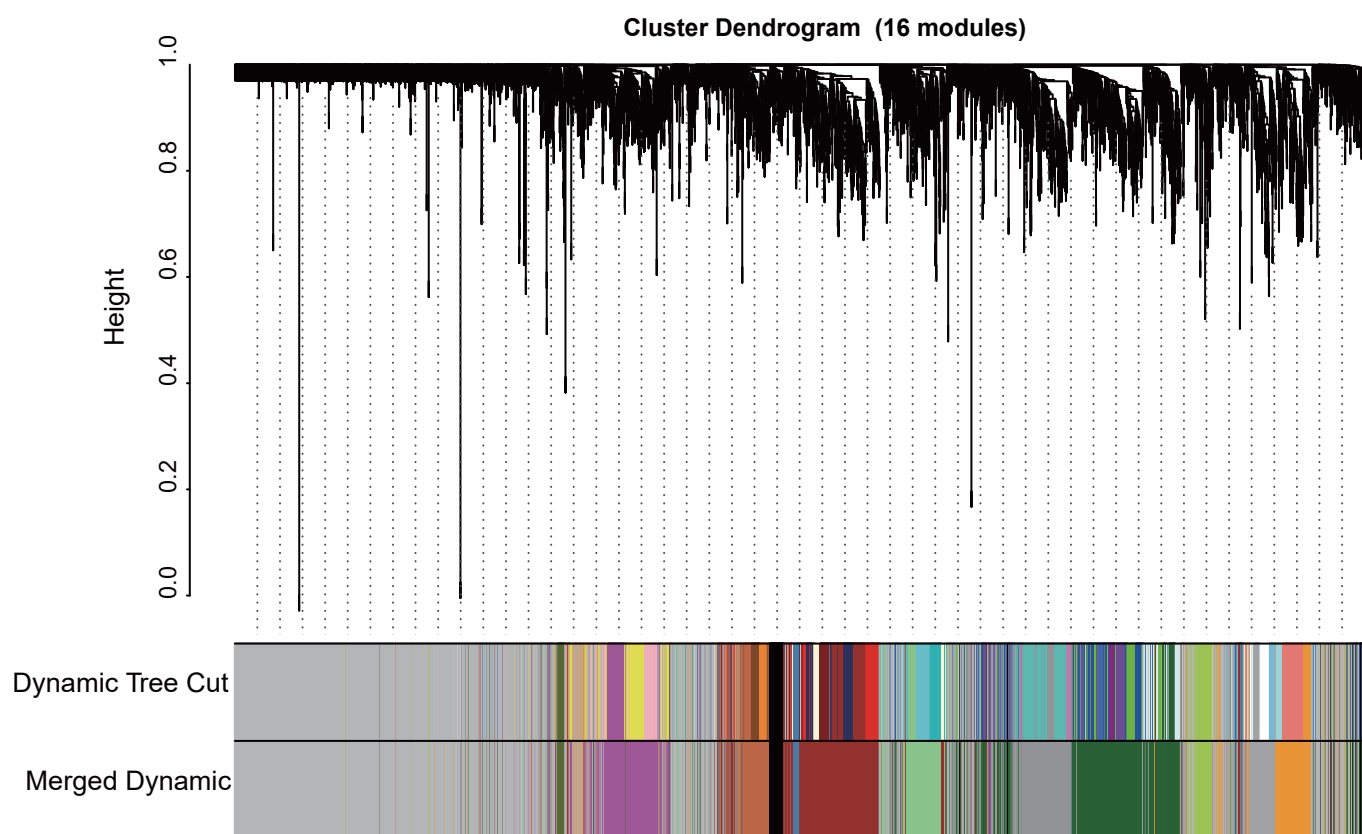
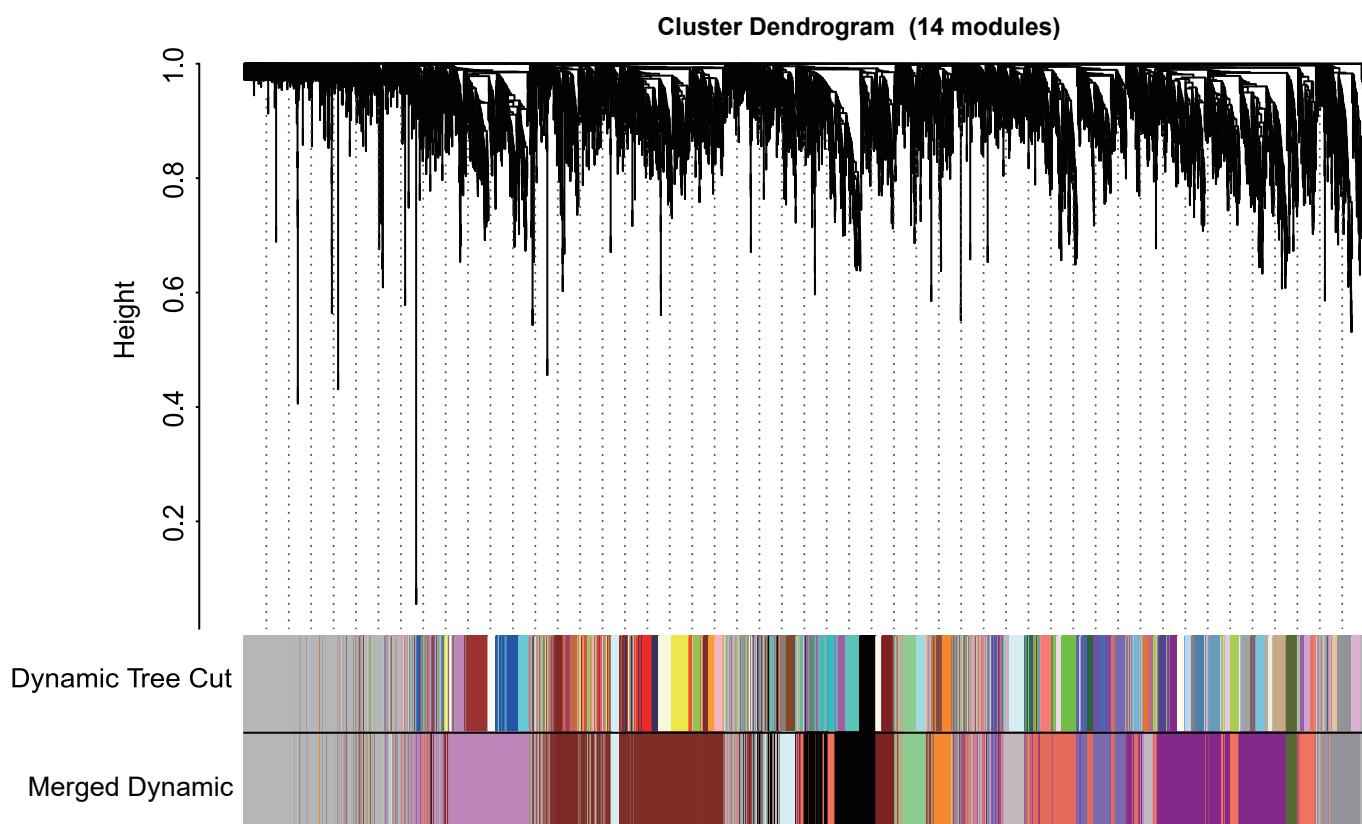


A**B**

Supplementary Figure S3: Distribution of DEGs in each module, with each branch representing one gene of the WGCNA result. WGCNA analysis uses the DEGs in leaves (A) and roots (B) of the three *Saccharum* species