

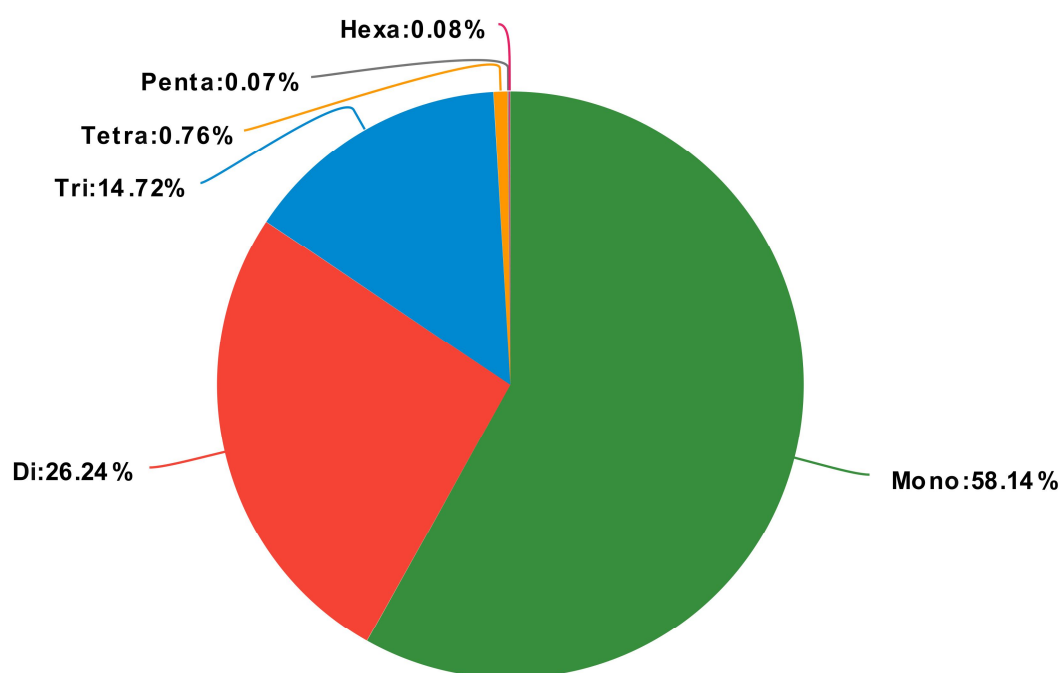
## *Supplementary Material*

### **Characterization of the de novo *Cinnamomum chago* (Lauraceae) transcriptome reveals candidate genes for terpenoid, fatty acid biosyntheses and abiotic stress**

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**Supplementary Figure 4** Frequency distribution of EST-SSRs based on motif types in *C. chago*