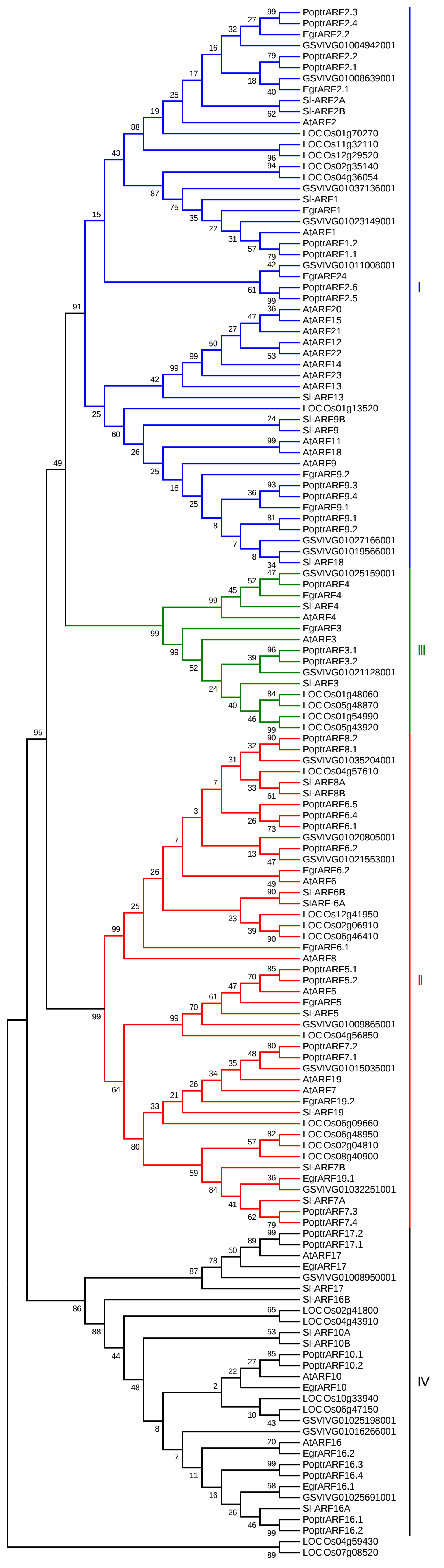
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I
II
III
IV

Supplementary Figure 5. Phylogenetic relationships between ARF proteins from woody species (poplar, *Eucalyptus* and grapevine) herbaceous species (*Arabidopsis*, tomato and rice). Full-length protein sequences were aligned by using the Clustal_X program. The phylogenetic tree was constructed by using the MEGA5 program and the neighbor-joining method with predicted full-length ARF proteins. Bootstrap support is indicated at each node.