

Genome-Wide Expression of WRKY Transcription Factors in Blueberry: Go Wild and Duplicate

Winder Felipez ^{1,2,†}, Jennifer Villavicencio ^{2,3,†}, Valeria Oliveira Nizolli ², Camila Pegoraro ², Luciano da Maia ² and Antonio Costa de Oliveira ^{2,*}

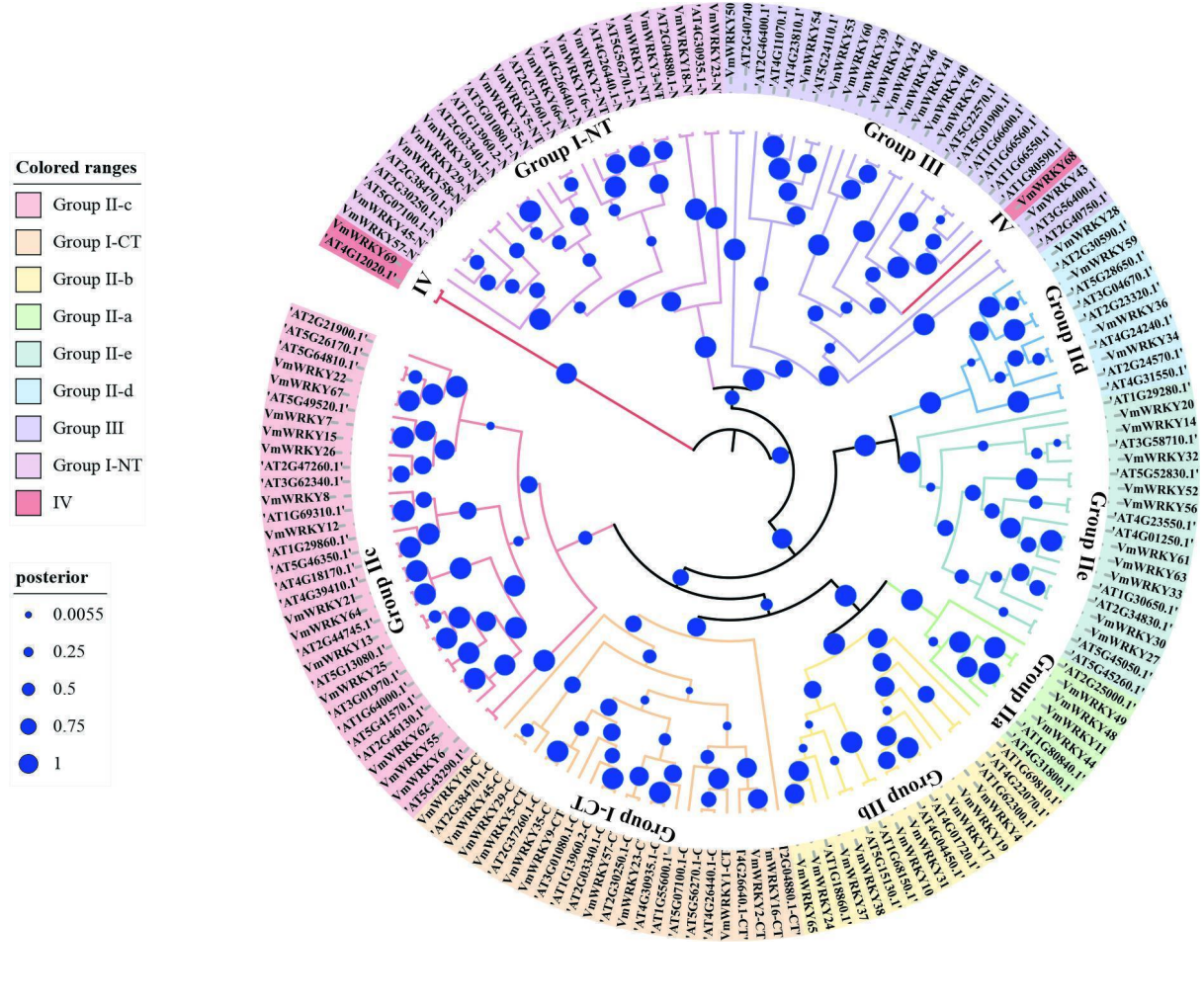


Figure S5: The unrooted phylogenetic tree was generated with the BEAST v.2.5 software, using the complete amino acid sequences of the 72 WRKY proteins from *Arabidopsis thaliana* genome and 69 WRKY proteins from *Vaccinium myrtillus* genome by Neighbor-Joining method, with the UPGMA clustering method and 1000 bootstrap repeats. WRKY protein subfamilies (I, IIa, IIb, IIc, IId, IIe and III).