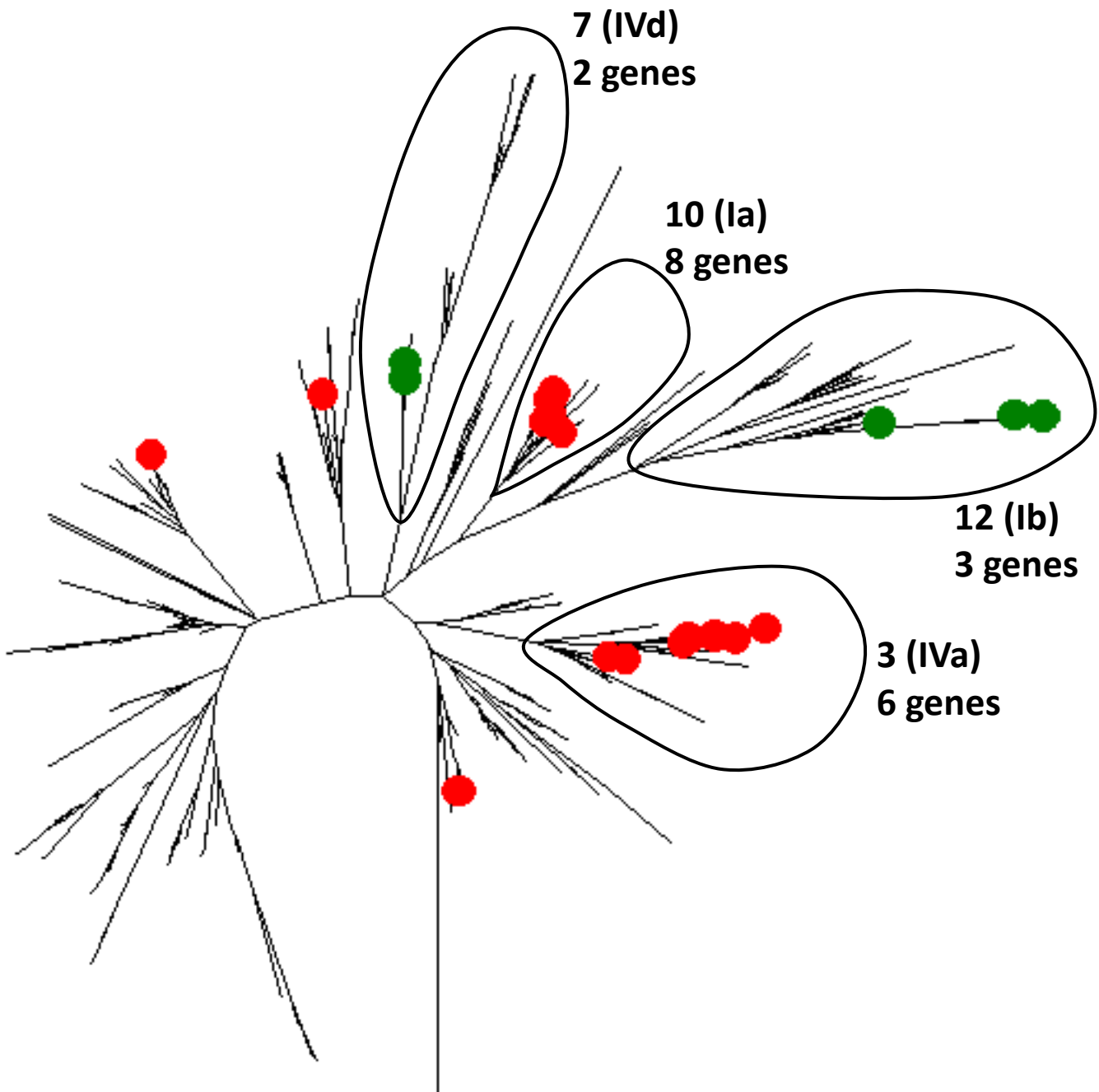


a.



b.

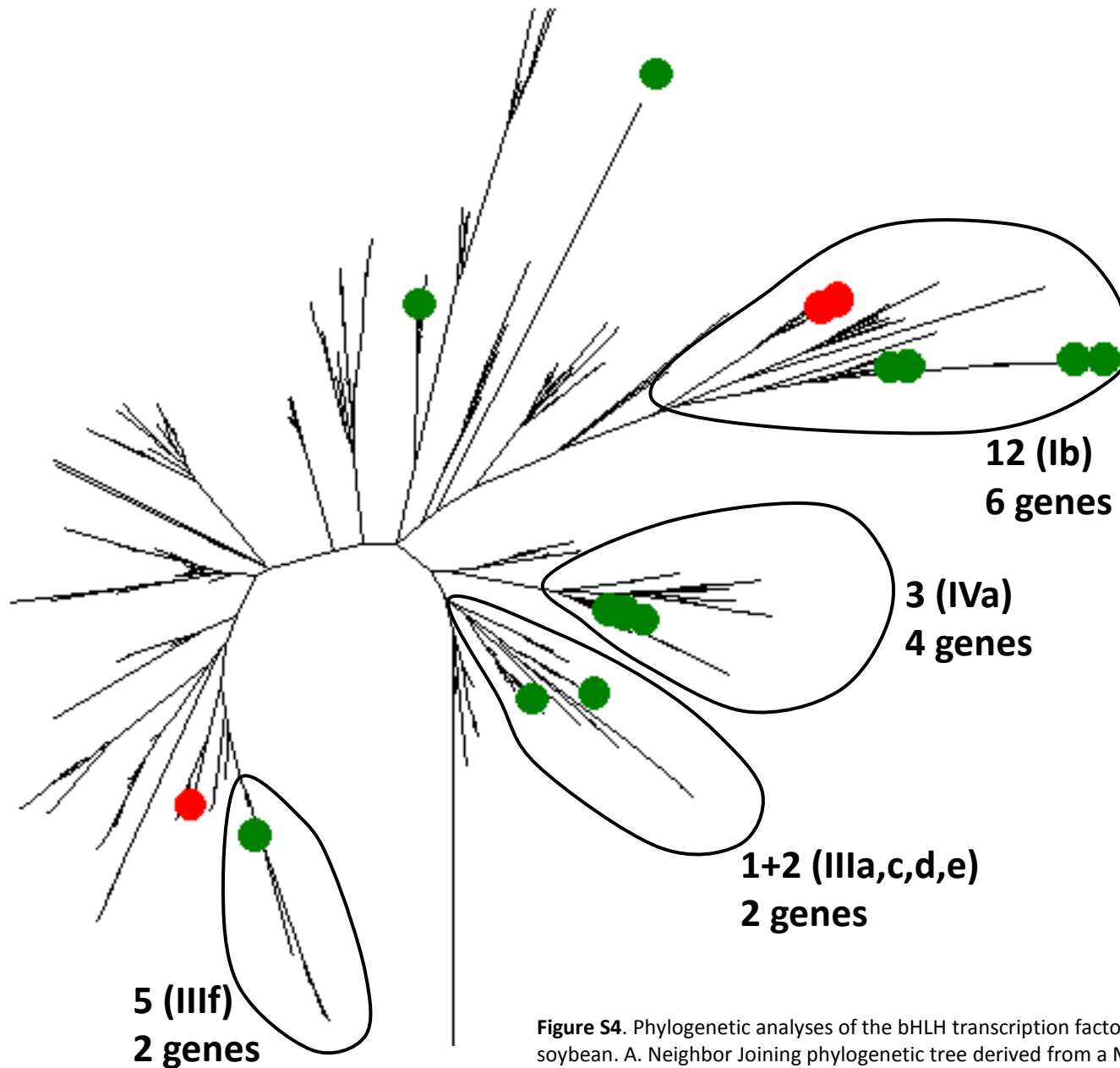


Figure S4. Phylogenetic analyses of the bHLH transcription factor family from soybean. A. Neighbor Joining phylogenetic tree derived from a MUSCLE alignment of the full length proteins. Red dots denote 8-fold down-regulation and green dots 8-fold up-regulation in roots. B. The same phylogenetic tree as shown in A. but with the inducibilities in leaf.