Supplementary data:

Genomewide identification and expression analysis of the ARF gene family in apple

Xiao-Cui Luo, Mei-Hong Sun, Rui-Rui Xu, Huai-Rui Shu, Jia-Wei Wang and Shi-Zhong Zhang

J. Genet. **93**, 785–797

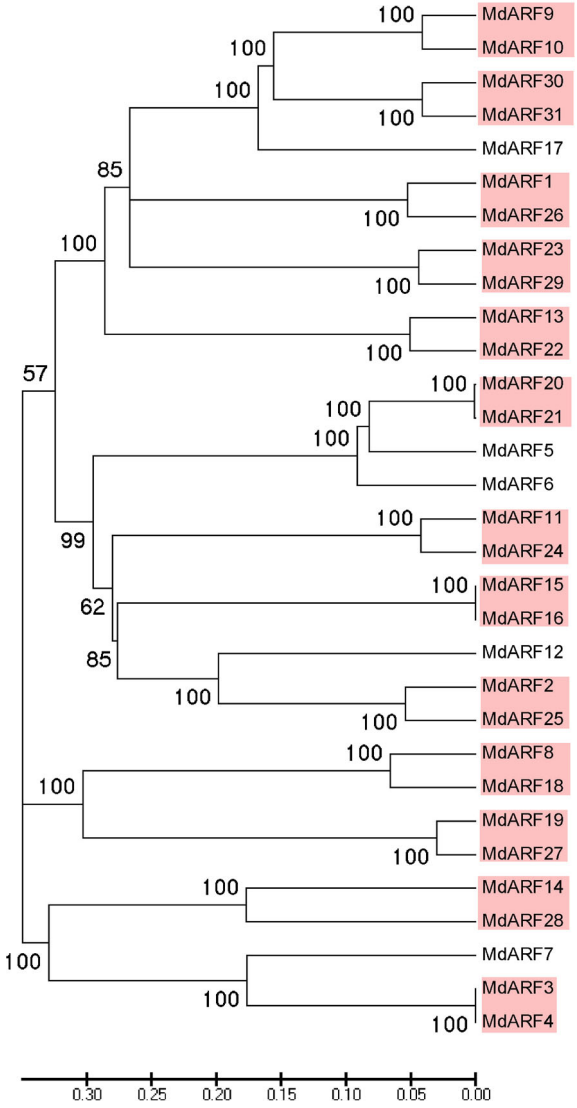

Figure 1. Phylogenetic relation of apple ARF genes. The phylogenetic tree was constructed based on a complete protein sequence alignment of *MdARFs* by the neighbour-joining method with bootstrapping analysis (1000 replicates). The scale bar represents 0.05 amino acid substitutions per site. Paralogous gene pairs are marked by a red box.