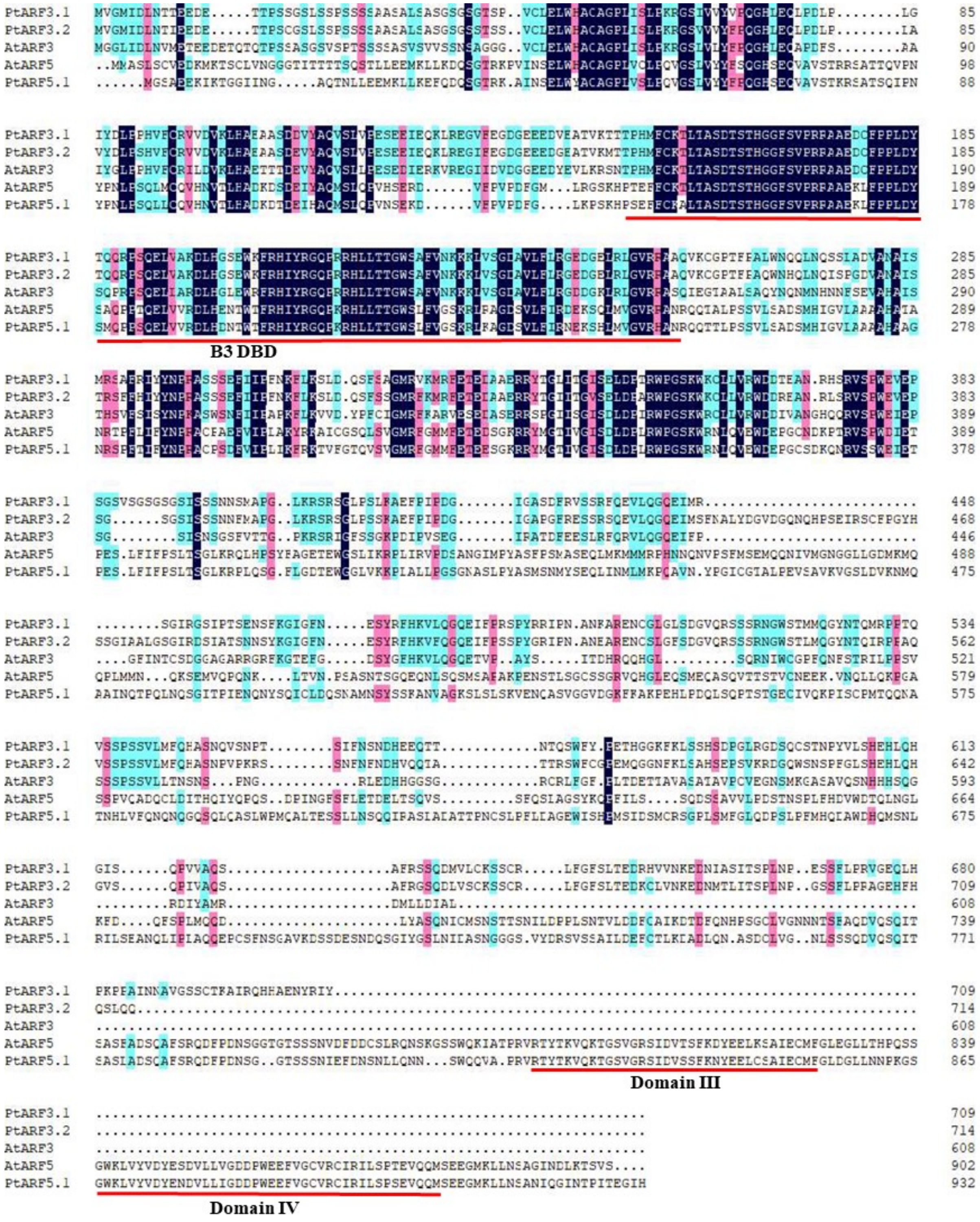

Fig. S1 Sequence alignment of PagARF3.1, PagARF3.2, PagARF5.1, Arabidopsis ARF3 and ARF5. PagARF3.1, PagARF3.2 and ARF3 are lack of the C-terminal dimerization domain (domain III and IV).