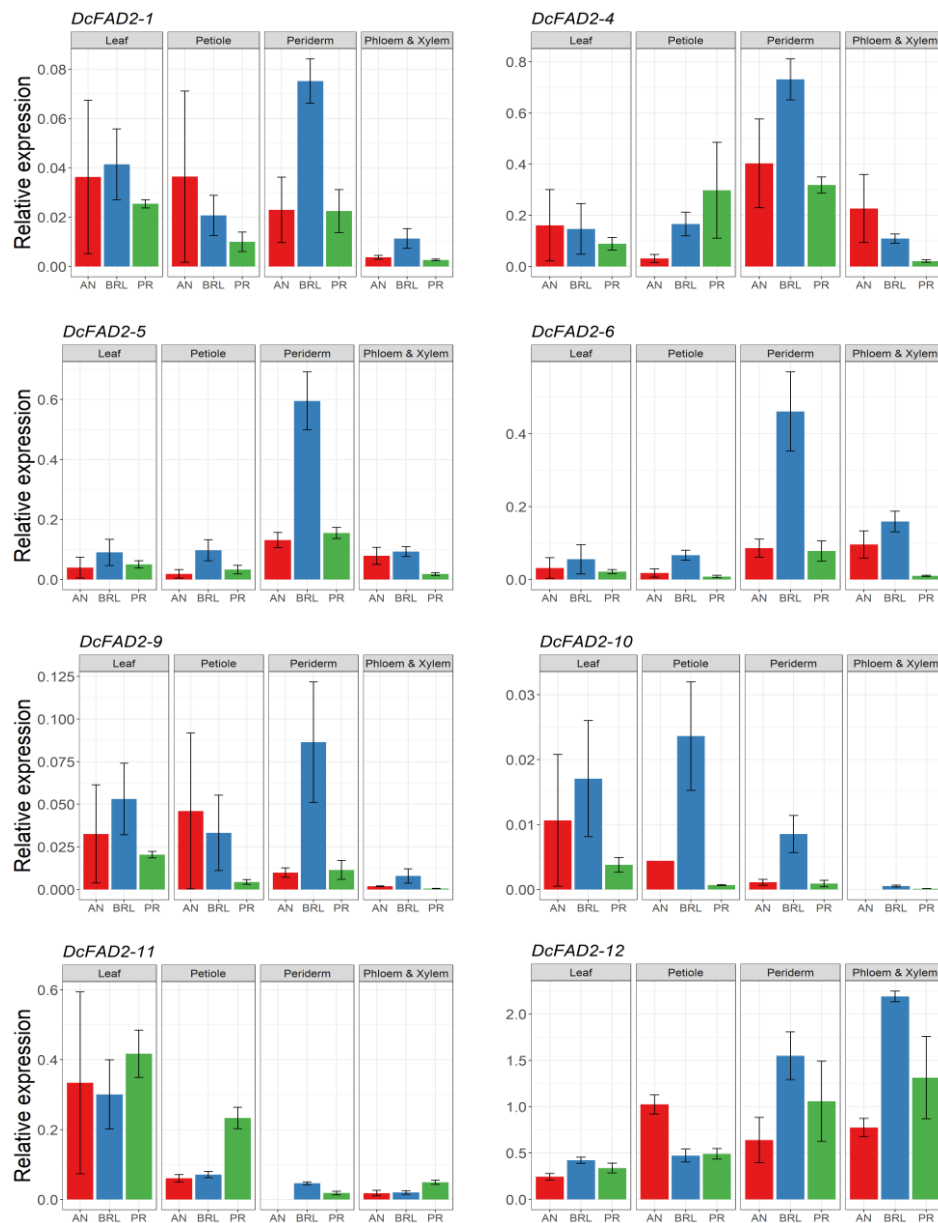




**Figure S9** Tissue-specific expression profiles of additional *FAD2* genes associated with minor QTLs (*DcFAD2-1*, *DcFAD2-4*, *DcFAD2-6*, *DcFAD2-9*, *DcFAD2-11*) or not associated with QTLs (*DcFAD2-5*, *DcFAD2-10*, *DcFAD2-12*). The levels of RNA transcripts were analysed in leaf, petiole, periderm, and a mixture of phloem and xylem of Anthonina (AN), Breeding line (BRL) and Presto (PR). Data represent means of three individuals of each cultivar with error bars indicating standard error.