

S5 Table. Identity (%) between MYCs transcription factors sequences of *Arabidopsis thaliana* and *Fragaria vesca* obtained by multiple alignment.

		<i>Fragaria vesca</i>	
		MYC2	MYC2-like
<i>Arabidopsis thaliana</i>	MYC2	58.51	41.54
	MYC3	54.16	38.88
	MYC4	58.27	38.92
	MYC5	45.56	38.69

Bold numbers indicate the highest identity of *F. vesca* MYCs comparing to *Arabidopsis*.