

Table S1. Predicted amino acid sequences for FvJAZ1/8

<i>Arabidopsis thaliana</i>			BLASTp				<i>Fragaria vesca</i>			
Protein ^a	Accession No.	Length (aa)	Score	Query cover (%)	E-value	Identity (%)	Predicted protein	Accession No.	Name ^b	Length (aa)
AtJAZ1	NP_564075	253	134	99	5e-38	38.7	TIFY 10A	XP_004287655	FvJAZ1	302
AtJAZ8	NP_564349	131	93.6	96	2e-25	46.2	TIFY5A	XP_004293626	FvJAZ8	130

^aAtJAZ1/8 amino acid sequences were used as queries and *Fragaria vesca* Refseq-protein database as subject.

^bThis refers to gene names assigned in the present research.