

**Table S3.** Expression analysis of all potential SI-candidate genes by SqRT-PCR.

SI candidate genes	Expression levels in various tissues								Expression levels in different stages after self-pollination and cross-pollination										
F library		B	L	P	F	Po	Sti	Sty	O		0h	4h	1d	2d	3d	4d	5d	6d	7d
H102	A1									C1									
	A2									C2									
H115	A1									C1									
	A2									C2									
H128	A1									C1									
	A2									C2									
H264	A1									C1									
	A2									C2									
H304	A1									C1									
	A2									C2									
H428	A1									C1									
	A2									C2									
H543	A1									C1									
	A2									C2									
H556	A1									C1									
	A2									C2									
H858	A1									C1									
	A2									C2									
H1092	A1									C1									
	A2									C2									
<b>R library</b>																			
F15	A1									C1									
	A2									C2									
F65	A1									C1									
	A2									C2									
F78	A1									C1									
	A2									C2									
F960	A1									C1									
	A2									C2									
F966	A1									C1									
	A2									C2									
F1166	A1									C1									
	A2									C2									
F250	A1									C1									
	A2									C2									
F465	A1									C1									
	A2									C2									
F499	A1									C1									
	A2									C2									
F519	A1									C1									
	A2									C2									
F546	A1									C1									
	A2									C2									

Table S3. Cont.

SI candidate genes	Expression levels in various tissues						Expression levels in different stages after self-pollination and cross-pollination							
R library														
F1019	A1							C1						
	A2							C2						
F580	A1							C1						
	A2							C2						
F685	A1							C1						
	A2							C2						
F822	A1							C1						
	A2							C2						

A1, “Wuzishatangju”; A2, “Shatangju”; C1, “Wuzishatangj” × “Wuzishatangju”; C2, “Wuzishatangju” × “Shatangju”; B, Buds; L, Leaves; P, Petals; F, Filaments; Po, Pollen; Sti, Stigmas; Sty, Styles; O, Ovaries.