







			Pollen donors			Controls	S-shared vs. Controls	Expressed in stigma ?
			F1					
S-haplotypes	1		<i>S12</i>	<i>S12</i>	<i>S12</i>			
	2		<i>S07</i>	<i>S02</i>	<i>S01</i>			
Plants			1	4	1			
Pollen receptors (stigma)		<i>S12</i>	<i>S07</i>	1	5/5	NA	NA	?
	F1	<i>S12</i>	<i>S02</i>	4	5/5	7/10		yes*
		<i>S12</i>	<i>S01</i>	1	5/5	4/5	 	?
Controls			NA	10/10	3/4			
S-shared vs. Controls			NA	 				
Expressed in pollen ?			?	?	yes*			

Figure S12 Summary of cross-pollinations realized for individuals from collection F0 and F1 having *S*-haplotype *S12* (*B17*-*B18*). See Figure S1 for legend details.

Haplotype *S12* was not typed in collection F1 because of no available specific primers for *B17* or *B18*. The presence of *S12* in collection F1 was thus deduced from the segregation pattern in offspring of individual 9 of (genotype *S03-S12*; see Figure 3 and Table 3). In collection F1, we thus considered that *S12* was present when *S03* (*B06*) was absent.