

S3 Table: Flowering time of transgenic plants over-expressing miR156-resistant *SPL4*, or point mutations in this gene.

Lines	Early flowering phenotype			# T1 plants
	Strong	Medium	Weak	
<i>35S::SPL4</i> ^a	39%	50%	11%	28
<i>35S::SPL4</i> Δ m1 ^b	0%	45%	55%	47
<i>35S::SPL4</i> Δ m2 ^c	16%	54%	30%	63
<i>35S::SPL4</i> Δ m3 ^d	4%	60%	37%	104

^a*SPL4* with no miR156 target site, expressed under the regulation of the CaMV 35S promoter.

^b*SPL4* Δ with the same point mutation as *spl4-1*

^c*SPL4* Δ with the same point mutation as *spl4-2*

^d*SPL4* Δ with the same point mutation as *spl4-3*