**Supplementary Table S1.** Nucleotide substitutions detected in the overexpression experiments. For each of the nine infiltrated plants, the substitutions observed in the clonal sequences analyzed at the overexpressed (AtCDA1 to AtCDA9) and control (pBIN61-infiltrated) regions are shown. In some cases, a given substitution is present in several clonal sequences from the same sample and the number of times it appears is indicated between parentheses. Nucleotide positions are given according to CaMV isolate W260, GenBank accession JF809616.1.

	Number			Number	
Sample	of clones	Mutation	Sample	of clones	Mutation
AtCDA1	106	A91G	pBIN61	106	T72C (2)
		G129A			A84G
		C131T			T98C
		C132T			C135G
		C143T			A162T (2)
		G148T			C174A
		G148A		_	G181T
		G153A			G181A (6)
		C157A (2)			C206T
		G159A			G210A
		C168T			A215G
		G169A			C224A
		C174A			A235G
		G175A			G237T
		G181A (27)			
		G190A (3)			
		C199T			
		G210A			
		C213T (3)			
		C213A			
		G214A (2)			
		A215G			
AtCDA2	106	T72C	pBIN61	106	T80C
		A77G			A161G
		A84G			C196A
		G126A			G221T
		T154C			A232G
		G169A			
		G181A (3)			
		C186G			
		A202C			
		T223C			
		G231A			
1.CD 12	2.4	A251G	DD1//1	2.4	1020
AtCDA3	24	A101G	pBIN61	24	A82G
LCD 11	2.4	C180T	DD1//1	2.4	C1 (OT
AtCDA4	24	C125T	pBIN61	24	G169T
1.CD 15	26	T70.C	DD1//1	26	T223C
AtCDA5	26	T79C	pBIN61	26	A121G
		C117T	á		A136G

		T140A			C186A C189T
1 65 16		C165G (3)	DD164		A212G
AtCDA6	25	C147A G224T	pBIN61	25	C135G
		022+1			T193C
AtCDA7	24	T96C	pBIN61	24	
		A103G			
		T140C			
		C186T			
AtCDA8	24	T114C	pBIN61	24	G68C
		C157T(5)			T98C
					G127C
					G214T
		T195C			T248C
AtCDA9	106	G85T	pBIN61	106	T114C
		C95T			T119C
		G126A			T128C
		T128C			C143T
		C146A (2)			C147A
		A161G			G153T
		A163G			T187G
		T176C			A212G
		G181A (2)			G214A
		G218C			G221T
		G231A			A224C
		A235G			
		C241T			