

**Table S1** Quantification of the centromere clustering defect in *corolla* mutants.

Genotype	Number of CID foci (%)						<i>n</i>	Average ± SD
	1	2	3	4	5	6		
<b>wild type</b>	16 (42.1)	13 (34.2)	9 (23.7)	0 (0)	0 (0)	0 (0)	38	1.80 ± 0.80
<b><i>corolla</i></b>	0 (0)	0 (0)	11 (34.4)	14 (43.8)	4 (12.5)	3 (9.4)	32	3.97 ± 0.93

The number of oocytes in region 3 of the germarium with a given number of CID foci are shown, followed in parentheses by the frequency that a given number of CID foci are seen in a genotype.