

Table S2. Mean time (in seconds \pm s.e.m.) separating meiosis II exit from NEBD of the male and female pronuclei and the interval between female and male NEBD in embryos of the indicated genotypes

Genotype	Meiosis II exit – male NEBD	Meiosis II exit – female NEBD	Δt (seconds)	<i>n</i>	<i>P</i> -value
<i>zyg-9(b244ts)</i>	555 \pm 12	630 \pm 14	73 \pm 8	9	–
<i>air-1(RNAi);zyg-9(b244ts)</i>	788 \pm 42	791 \pm 42	2.5 \pm 3	8	<0.001
<i>spat-1(RNAi)zyg9(b244ts)</i>	675 \pm 50	925 \pm 77	250 \pm 37	6	<0.001
<i>plk-1(RNAi);zyg-9(b244ts)</i>	866 \pm 55	1173 \pm 91	301 \pm 42	6	0.003

Data are represented in Fig. 2B. The number of embryos analysed (*n*) and *P*-values [calculated for Δt by two-tailed *t*-test comparing double mutants with the single *zyg-9(b244ts)*] are indicated.