

Table S1A

Genotype	Hatching Rate
<i>y w</i>	673/859 (78.3%)
<i>kumo<sup>M41-13</sup>/TM3</i>	738/912 (80.9%)
<i>kumo<sup>M41-13</sup></i>	0/779 (0%)
<i>w<sup>-</sup>;Myc-Kumo/nosgal4;kumo<sup>M41-13</sup></i>	693/917 (75.5%)
<i>w<sup>-</sup>;Myc-KumoNT/nosgal4;kumo<sup>M41-13</sup></i>	0/478 (0%)
<i>w<sup>-</sup>;Myc-KumoCT/nosgal4;kumo<sup>M41-13</sup></i>	287/622 (46.14%)

Table S1B

Genotype	Normal appendages
<i>y w</i>	663/712 (93.1%)
<i>kumo<sup>M41-13</sup>/TM3</i>	651/689 (94.4%)
<i>kumo<sup>M41-13</sup></i>	268/636 (42.1%)
<i>w<sup>-</sup>;Myc-Kumo/nosgal4;kumo<sup>M41-13</sup></i>	475/540 (87.9%)

Table S1A: Hatching rate of eggs from wild type, *kumo* mutant, *kumo* mutants expressing Kumo-NT, and Kumo-CT transgenes mothers.

Table S1B: Comparison of number of eggs having normal appendages from wild type, *kumo* mutant, and *kumo* mutants expressing full length kumo transgene mothers