

Genotype	Ave # progeny $\pm$ std error	% change	N=
Experiment 1:			
wild type	233 $\pm$ 62	--	8
<i>sma-2(e502)</i>	67 $\pm$ 57	-71%	8
<i>sma-9(qc3)</i>	156 $\pm$ 34	-33%	8
<i>sma-9(wk55)</i>	99 $\pm$ 61	-58%	8
<i>daf-4(e1364)</i>	33 $\pm$ 14	-86%	8
Experiment 2:			
wild type	263 $\pm$ 19	--	10
<i>sma-2(e502)</i>	77 $\pm$ 55	-71%	10
<i>sma-9(qc3)</i>	248 $\pm$ 34	-6%	10
<i>sma-9(wk55)</i>	162 $\pm$ 44	-38%	10
Experiment 3:			
wild type	243 $\pm$ 62	--	20
<i>sma-2(e502)</i>	122 $\pm$ 65	-50%	30
<i>sma-9(qc3)</i>	200 $\pm$ 38	-18%	26
<i>sma-9(wk55)</i>	153 $\pm$ 34	-37%	20
<i>daf-4(e1364)</i>	74 $\pm$ 31	-70%	26
Experiment 4:			
wild type	326 $\pm$ 37	--	30
<i>sma-2(e502)</i>	114 $\pm$ 36	-65%	30
<i>sma-9(wk55)</i>	218 $\pm$ 33	-33%	30
Experiment 4:			
wild type	243 $\pm$ 21	--	20
<i>daf-4(e1364)</i>	74 $\pm$ 31	-70%	26