

Genotype	mean RS± std. error	% change	P-value	N=
Life span				
wild type	17.6 ±0.6	--	--	72
<i>sma-2(e502)</i>	18.4 ±0.3	+5%	0.52	96
<i>daf-7(e1372)</i>	21.3 ±0.8	+21%	0.0001	97
<i>daf-2(e1370)</i>	36.2 ±1.6	+106%	<0.0001	72
<i>eat-2(ad465)</i>	21.3 ±0.9	+21%	0.0011	72
Reproductive span (mated)				
wild type	6.6 ±0.4*	--	--	15
<i>sma-2(e502)</i>	10.0 ±0.6*	+52%	0.0034	29
<i>daf-7(e1372)</i>	7.8 ±0.2	+18%	0.04	39
<i>daf-2(e1370)</i>	11.0 ±0.3*	+67%	<0.0001	55
<i>eat-2(ad465)</i>	12.0 ±0.4*	+82%	<0.0001	69

*Corrected for matricide (animals bagged at the last day were assumed to have ceased its reproduction on that day to calculate the mean RS, to avoid underestimation due to high matricide rate). Such correction calculates a value closer to the true mean RS, and is important when comparing the relative increases in reproductive span.