

Genotype	mean LS/RS± std. error	% change	P-value	N
Life span experiment 1:				
wild type	15.8 ±0.4	--	--	72
<i>sma-2(e502)</i>	18.2 ±0.7	+15%	0.0036	73
<i>daf-7(e1372)</i>	21.1 ±0.6	+34%	<0.0001	73
Life span experiment 2:				
wild type	12.7 ±0.5	--	--	69
<i>daf-2(e1370)</i>	40.7±2.5	+220%	<0.0001	70
<i>eat-2(ad465)</i>	19.3 ±1.0	+52%	<0.0001	69
Reproductive span experiment 1:				
wild type	3.9 ±0.3	--	--	12
<i>sma-2(e502)</i>	8.7 ±0.8*	+123%	<0.0001	13
<i>daf-7(e1372)</i>	5.6 ±0.2	+44%	<0.0001	17
Reproductive span experiment 2:				
wild type	3.7 ±0.2*	--	--	30
<i>daf-2(e1370)</i>	5.9 ±0.3	+59%	<0.0001	45
<i>eat-2(ad465)</i>	8.2 ±0.4	+122%	<0.0001	44

*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.