

S1 Table. Starvation survival analysis of Figure 1A.

Genotype	Half-life	SEM	p-value (vs. wild type)	p-value (vs. <i>daf-16</i>)	R ²
wild type	10.5	0.58	N/A	3.8x10⁻⁶	0.966
<i>daf-16</i>	4.7	1.01	3.8x10⁻⁶	N/A	0.996
<i>daf-16; Pdaf-16::DAF-16</i>	10.8	1.15	0.681	6.7x10⁻⁵	0.961
<i>daf-16; Pges-1::DAF-16</i>	9.1	0.76	0.013	5.0x10⁻⁵	0.972
<i>daf-16; Punc-119::DAF-16</i>	7.2	1.34	0.002	0.014	0.986
<i>daf-16; Pcol-12::DAF-16</i>	8.6	1.26	0.025	0.003	0.987