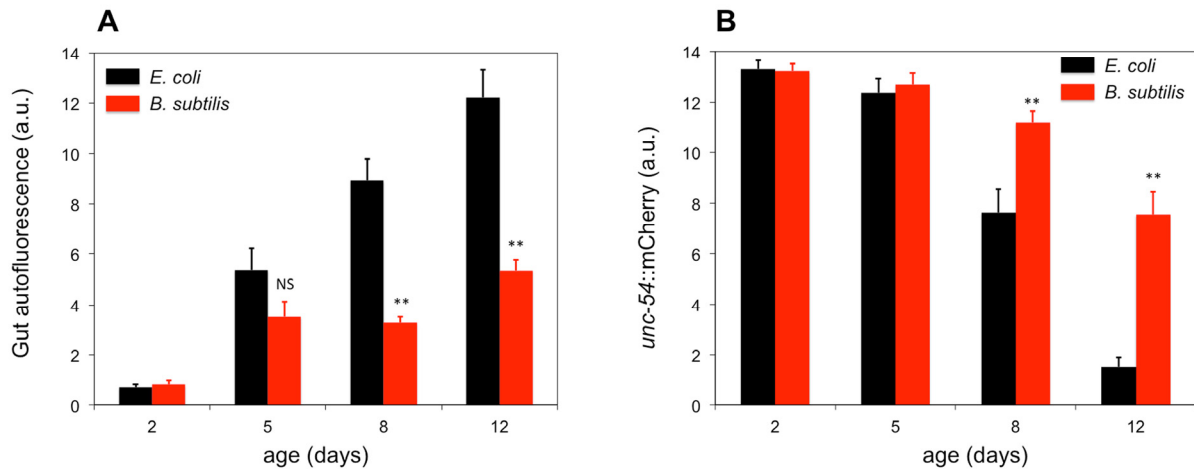
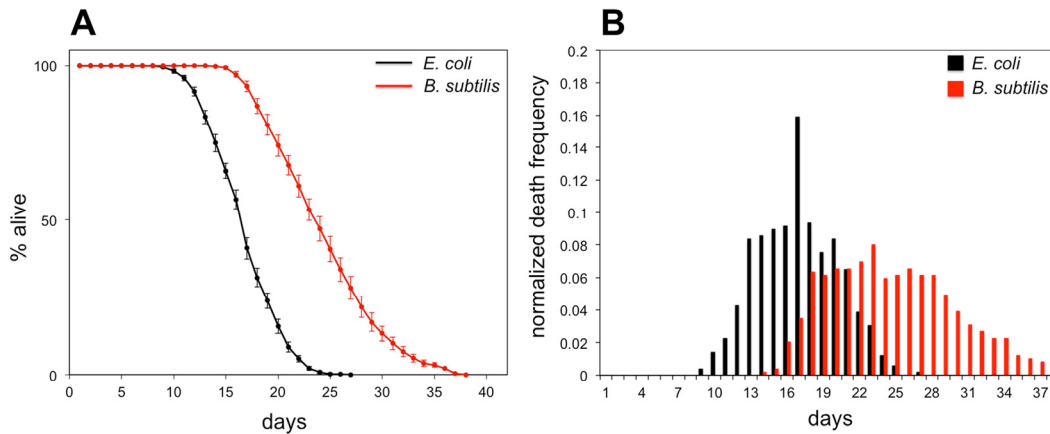


SUPPLEMENTAL DATA

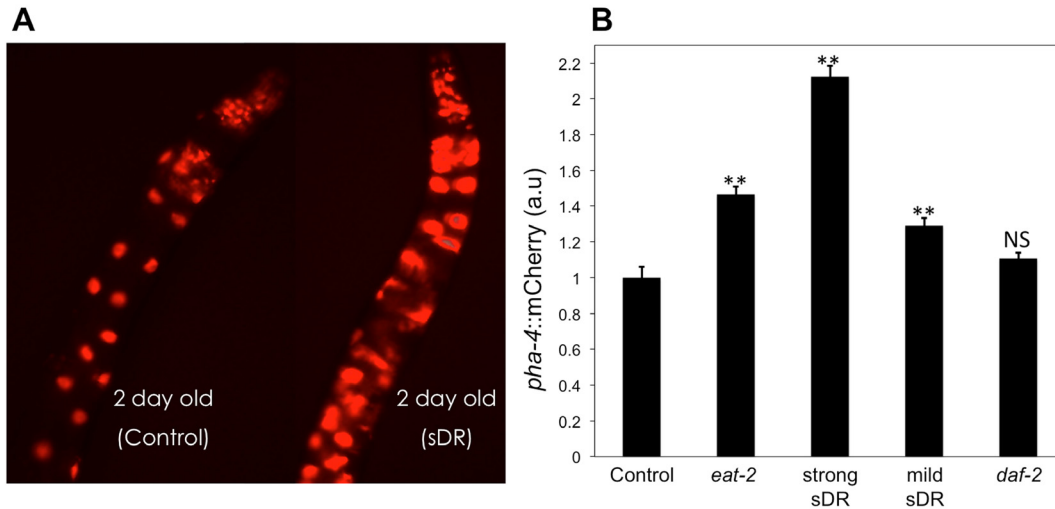


**Figure S1. Gut autofluorescence and *unc-54::mCherry* reporter as indicators of the overall health of *B. subtilis* and *E. coli* fed worms.** (A) Gut autofluorescence intensity for *B. subtilis* and *E. coli* fed adult hermaphrodites during aging. y-axis shows levels of gut autofluorescence in arbitrary units. x-axis shows age of worms. (B) *unc-54::H1::mCherry* fluorescent marker expression for *B. subtilis* and *E. coli* fed adult hermaphrodites during aging. y-axis shows levels of fluorescent expression in arbitrary units. x-axis shows age of worms. (A-B) Bars indicate the mean fluorescent marker expression  $\pm$  S.E.M. n = 10-15 for each group (\*\*p < 0.01; NS p > 0.05; Student's t test).

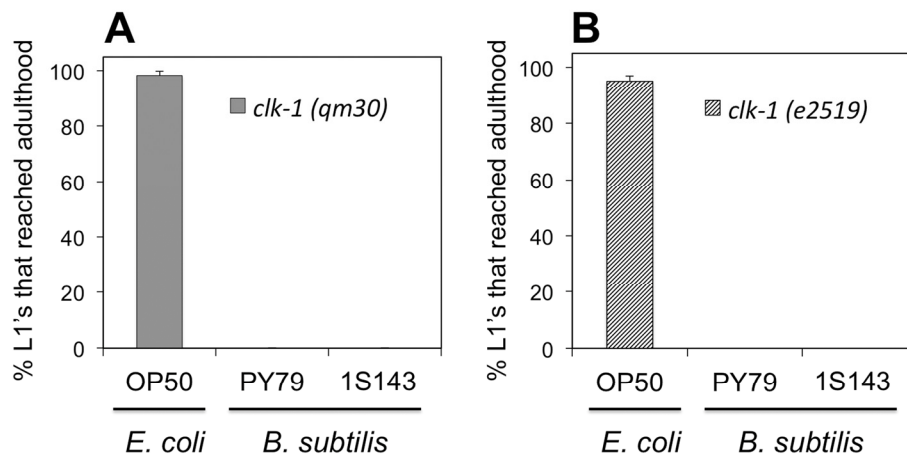


	<i>E. coli</i> (OP50)	<i>B. subtilis</i> (PY79)
Number of worms	491	488
Median life span ( $\pm$ SD)	16.4 $\pm$ 0.6	23.7 $\pm$ 1.7
Life span difference	44.5%	
P. value (log rank)	<0.0001	

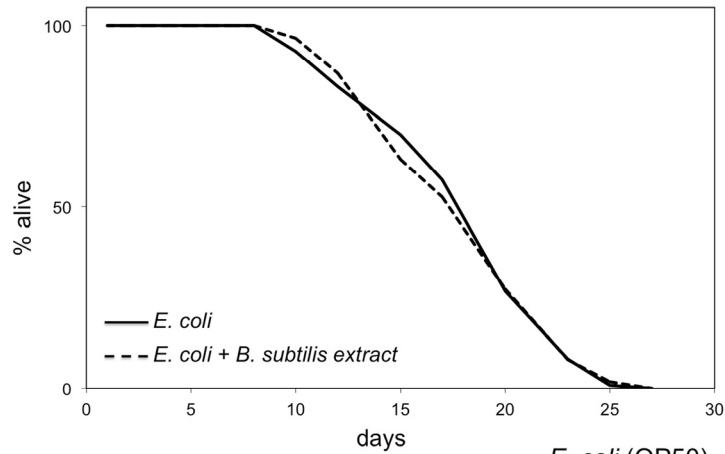
**Figure S2. *B. subtilis* fed worms live longer than *E. coli* fed worms and display different demographics of death.** (A) Represented are the life span curves for adult hermaphrodite worms maintained on *E. coli* (OP50) or on *B. subtilis* (PY79). The number of dead worms was counted on a daily basis. The worm populations were divided into 10 plates. The points in the life span curves represent the daily mean percentage of alive worms in the 10 plates  $\pm$  S.E.M. y-axis indicates percentage of worms that are alive. x-axis indicates day of adulthood. (B) Represented are the daily death frequencies normalized to their respective sample size.



**Figure S3. *pha-4::mCherry* reporter as a worm nutritional indicator.** (A) Representative pictures of *pha-4::H1::mCherry* fluorescent marker expression for 2 day old WT adult hermaphrodites, which were exposed to plenty of food (Control) or dietary restricted (sDR). (B) *pha-4::H1::mCherry* fluorescent marker expression for 2 day old WT adult hermaphrodites, which were exposed to plenty of food (Control) or dietary restricted (sDR); and for 2 day old *eat-2(ad1116)* and *daf-2(e1370)* mutant adult hermaphrodites, which were exposed to plenty of food. For details on dietary restriction conditions see the Materials and Methods section. y-axis shows levels of fluorescent expression in arbitrary units. x-axis shows the type of worms and the diet condition. Bars indicate the mean fluorescent marker expression  $\pm$  S.E.M.  $n = 15$  for each group (\*\* $p < 0.01$ , Student's t test). (A-B) In all cases worms were fed *E. coli*.

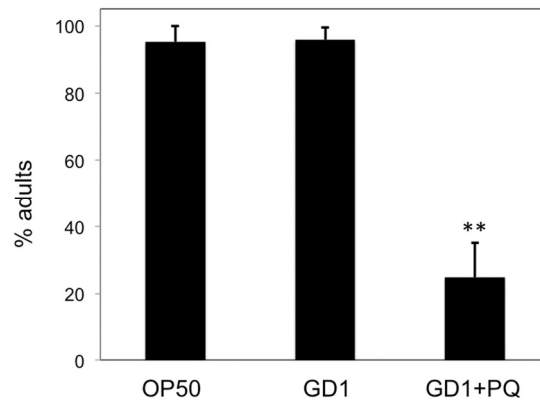


**Figure S4. *clk-1* mutant worms are unable to develop into adults when fed the *B. subtilis* diet.** (A) Percentage of *clk-1(qm30)* L1 larvae that developed into adults after 3.9 days of feeding on *E. coli* (OP50), *B. subtilis* (PY79) or the spore-less *B. subtilis* (1S143). (B) Percentage of *clk-1(e2519)* L1 larvae that developed into adults after 4.0 days of feeding on *E. coli* (OP50), *B. subtilis* (PY79) or the spore-less *B. subtilis* (1S143). (A-B) y-axis shows percentage of L1 larvae that reached adulthood. x-axis shows the diet used. Bars indicate the mean value  $\pm$  S.E.M.  $n = 140-160$  worms per group.

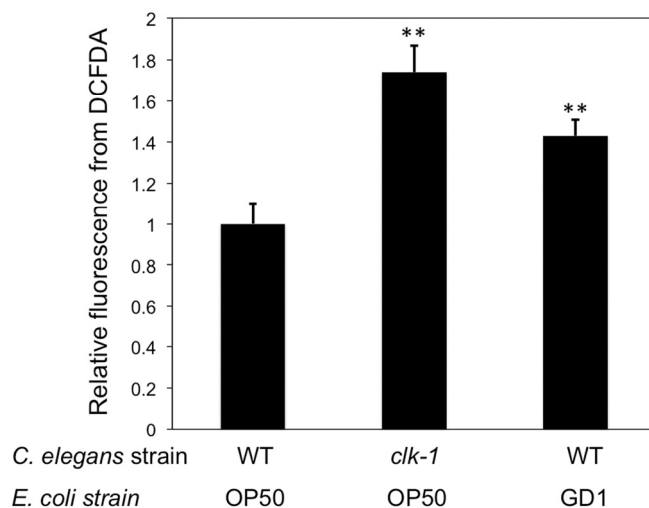


	<u>E. coli (OP50)</u>	<u>E. coli (OP50) + B. subtilis extract</u>
Number of worms	113	114
Median life span ( $\pm$ SD)	17.4 $\pm$ 0.6	17.1 $\pm$ 1.1
Life span difference	-1.4%	
P. value (log rank)	>0.05	

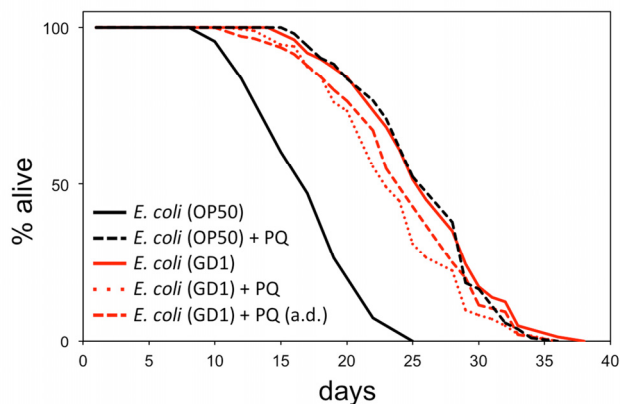
**Figure S5. *E. coli* fed worms supplemented with coQ-less *B. subtilis* extract do not alter their longevity.** Represented are the life span curves for adult hermaphrodite worms maintained on *E. coli* with or without supplementation of coQ-less *B. subtilis* extract. The spore-less *B. subtilis* (1S143) strain was used to prepare the *B. subtilis* extract with the purpose of avoiding the presence of *B. subtilis* spores in the extract. y-axis indicates percentage of worms that are alive. x-axis indicates day of adulthood.



**Figure S6. Treatment with a mild dose of paraquat severely affects the development of worms feeding on coQ-defective *E. coli* (GD1) bacteria.** Development of synchronized wild type N2 L1 larvae after 3 days feeding on *E. coli* OP50 and *E. coli* GD1 with or without PQ treatment (0.1 mM). y-axis shows percentage of individuals that reached adulthood after 3 days. x-axis shows the type of *E. coli* and the treatment. Bars indicate the mean value  $\pm$  S.D. n = 114-127 worms per group. \*\*p < 0.001, Student's t test.



**Figure S7. *E. coli* (GD1) fed wild type worms and *E. coli* (OP50) fed *clk-1(qm30)* worms have higher ROS levels than *E. coli* fed WT worms.** Bars indicate the relative mean fluorescent marker expression  $\pm$  S.E.M difference relative to the *E. coli* fed WT worms.  $n = 14-20$  for each group (\*\* $p < 0.01$ , Student's  $t$  test). y-axis indicates relative fluorescence from DCFDA. x-axis indicates diet and type of worms.



	OP50	OP50+PQ	GD1	GD1+PQ	GD1+PQ(a.d.)
Number of worms	68	103	208	140	184
Median life span ( $\pm$ SD)	16.5 $\pm$ 1.0	25.5 $\pm$ 1.1	25.2 $\pm$ 1.1	22.9 $\pm$ 0.9	23.8 $\pm$ 1.8
Life span difference		54.5%	52.7%	38.8%	44.2%
P. Value (log rank)		< 0.0001	< 0.0001	< 0.0001	< 0.0001

**Figure S8. Life span extension of paraquat-treated worms depends on the presence of coQ in the *E. coli* diet.** Represented are the life span curves for adult worms maintained on coQ-active *E. coli* OP50 or coQ-deficient *E. coli* GD1 with or without PQ treatment (0.1 mM). Worms were subjected to PQ treatment since L1, or since adulthood (a.d.). y-axis indicates percentage of worms that are alive. x-axis indicates day of adulthood.

**Table S1. Supporting life span data for Figure 1**

<i>C. elegans</i> strain	Bacterial food strain	Sample size	Median life span ± SD	% Life span change	P. value (log rank)
Wild type (N2)	<i>E. coli</i> (OP50)	178	14.2±0.8		
Wild type (N2)	<i>B. subtilis</i> (168)	169	20.4±0.5	+43.1%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (3610)	148	22.2±1.5	+56.0%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (PY79)	170	22.6±1.0	+58.5%	<0.0001

<i>C. elegans</i> strain	Bacterial food strain	Sample size	Median life span ± SD	% Life span change	P. value (log rank)
Wild type (N2)	<i>E. coli</i> (OP50)	107	15.5±0.3		
Wild type (N2)	<i>B. subtilis</i> (PY79)	118	25.2±0.4	+62.6%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (1S143)	94	22.7±1.2	+46.5%	<0.0001

<i>C. elegans</i> strain	Bacterial food strain	Sample size	Median life span ± SD	% Life span change	P. value (log rank)
Wild type (N2)	<i>E. coli</i> (OP50)	76	15.9±0.4		
Wild type (N2)	<i>B. subtilis</i> (PY79)	132	23.0±0.7	+44.7%	<0.0001
Wild type (N2)	UV killed <i>B. subtilis</i> (PY79)	81	22.7±0.8	+42.8%	<0.0001
Wild type (N2)	UV killed <i>E. coli</i> (OP50)	96	19.2±0.4	+20.8%	<0.001

<i>C. elegans</i> strain	Bacterial food strain	Sample size	Median life span ± SD	% Life span change	P. value (log rank)
Wild type (N2)	<i>E. coli</i> (OP50)	107	15.5±0.3		
Wild type (N2)	<i>B. subtilis</i> (PY79)	118	25.2±0.4	+62.6%	<0.0001
Wild type (N2)	Worms developed from L1 to late L4 on <i>E. coli</i> (OP50), then switched to <i>B. subtilis</i> (PY79)	111	19.3±2.4	+24.5%	<0.0001

**Table S2. Nutritional composition of *E. coli* vs. *B. subtilis***

	<i>E. coli</i> (OP50)	<i>B. subtilis</i> (PY79)
% Water content	78.02	78.75
Protein *	84.72	77.04
Fat *	0.41	0.28
Carbohydrate *	4.51	9.79
Ashes *	10.37	12.89
Calories **	360.57	349.82

\* Grams per 100 g of dry bacteria

\*\* Calories per 100 g of dry bacteria

**Table S3. Development of *clk-1* mutant L1 larvae after 7 days on the *B. subtilis* diet with *E. coli* extract supplementation**

<b><i>C. elegans</i> strain</b>	<b>Bacterial food strain</b>	<b>Extract supplemented*</b>	<b>Extract dilution</b>	<b>Outcome upon extract supplementation</b>
<i>clk-1(qm30)</i>	<i>B. subtilis</i> (PY79)	<i>E. coli</i> (OP50)	Undiluted	L1s developed into adults and these produced progeny that arrested at L1-L2 stages
<i>clk-1(qm30)</i>	<i>B. subtilis</i> (PY79)	<i>E. coli</i> (OP50)	1:2	L1s developed into adults and some of these produced progeny that arrested at L1-L2 stages
<i>clk-1(qm30)</i>	<i>B. subtilis</i> (PY79)	<i>E. coli</i> (OP50)	1:5	L1s developed into adults and these laid dead embryos
<i>clk-1(qm30)</i>	<i>B. subtilis</i> (PY79)	<i>E. coli</i> (OP50)	1:10	Some L1s developed into adults
<i>clk-1(qm30)</i>	<i>B. subtilis</i> (PY79)	<i>E. coli</i> (OP50)	1:20	Most L1s developed into L4s

\* single dose of *E. coli* extract was used

**Table S4. Supporting life span data for Figure 3**

<b><i>C. elegans</i> strain</b>	<b>Bacterial food strain</b>	<b>Sample size</b>	<b>Median life span ± SD</b>	<b>% Life span change</b>	<b>P. value (log rank)</b>
Wild type (N2)	<i>E. coli</i> (OP50)	236	16.4±1.1		
Wild type (N2)	<i>E. coli</i> (OP50) + <i>E. coli</i> (OP50) extract	118	17.3±0.4	+5.5%	>0.05
Wild type (N2)	<i>E. coli</i> (OP50) + <i>E. coli</i> (OP50) extract (1:5 dilution)	188	16.7±0.8	+1.8%	>0.05
Wild type (N2)	<i>B. subtilis</i> (PY79)	247	24.0±1.0	+46.3%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (PY79) + <i>E. coli</i> (OP50) extract	117	18.4±1.0	+12.2%	<0.005
Wild type (N2)	<i>B. subtilis</i> (PY79) + <i>E. coli</i> (OP50) extract (1:5 dilution)	187	20.6±1.2	+25.6%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (PY79) + <i>E. coli</i> (OP50) extract (1:10 dilution)	79	20.7±1.4	+26.2%	<0.0001
Wild type (N2)	<i>E. coli</i> (OP50)	120	16.4±0.8		
Wild type (N2)	<i>E. coli</i> (OP50) + <i>E. coli</i> (GD1) extract	84	16.7±0.3	+1.8%	>0.05
Wild type (N2)	<i>E. coli</i> (OP50) + <i>E. coli</i> (GD1) extract (1:5 dilution)	108	16.4±0.4	+0.2%	>0.05
Wild type (N2)	<i>B. subtilis</i> (PY79)	92	24.1±0.9	+47.0%	<0.0001

Wild type (N2)	<i>B. subtilis</i> (PY79) + <i>E. coli</i> (GD1) extract	80	22.4±0.5	+36.6%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (PY79) + <i>E. coli</i> (GD1) extract (1:5 dilution)	113	24.1±0.6	+47.0%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (PY79) + <i>E. coli</i> (GD1) extract (1:10 dilution)	122	24.1±1.0	+47.0%	<0.0001

**Table S5. Supporting life span data for Figure 5**

<i>C. elegans</i> strain	Bacterial food strain	Sample size	Median life span ± SD	% Life span change	P. value (log rank)
Wild type (N2)	<i>E. coli</i> (OP50)	202	17.4±0.5		
Wild type (N2)	<i>E. coli</i> (OP50) + 10mM NAC	195	17.5±1.0	+0.6%	>0.05
Wild type (N2)	<i>B. subtilis</i> (PY79)	235	24.5±0.6	+40.8%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (PY79) + 10 mM NAC	102	22.1±1.0	+27.0%	<0.001

<i>C. elegans</i> strain	Bacterial food strain	Sample size	Median life span ± SD	% Life span change	P. value (log rank)
Wild type (N2)	<i>E. coli</i> (OP50)	150	17.2±0.6		
Wild type (N2)	<i>E. coli</i> (OP50) + 0.1 mM PQ	146	25.1±1.5	+45.9%	<0.0001
Wild type (N2)	<i>E. coli</i> (OP50) + 0.1 mM PQ (after development)	180	23.6±1.6	+37.2%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (PY79)	169	23.2±1.2	+34.9%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (PY79) + 0.1 mM PQ	151	27.2±2.0	+58.1%	<0.0001
Wild type (N2)	<i>B. subtilis</i> (PY79) + 0.1 mM PQ (after development)	180	25.1±1.8	+45.9%	<0.0001

**Table S6. Supporting life span data for Figure 6**

<i>C. elegans</i> strain	Molecular pathway or process affected	Bacterial food strain	Sample size	Median life span $\pm$ SEM	Bacterial food strain	Sample size	Median life span $\pm$ SEM	% Life span change	P. value (log rank)
Wild type (N2)		<i>E. coli</i> (OP50)	512	16.0 $\pm$ 0.3	<i>B. subtilis</i> (PY79)	465	23.2 $\pm$ 0.5	+45.1%	<0.0001
<i>daf-2(e1370)</i>	Insulin-like pathway	<i>E. coli</i> (OP50)	159	41.9 $\pm$ 1.1	<i>B. subtilis</i> (PY79)	90	48.0 $\pm$ 2.8	+14.5%	<0.001
<i>daf-16(mu86)</i>	Insulin-like pathway	<i>E. coli</i> (OP50)	346	13.9 $\pm$ 0.3	<i>B. subtilis</i> (PY79)	229	15.7 $\pm$ 0.5	+13.2%	<0.0005
<i>daf-16(mgDf50)</i>	Insulin-like pathway	<i>E. coli</i> (OP50)	117	11.7 $\pm$ 0.1	<i>B. subtilis</i> (PY79)	115	14.9 $\pm$ 0.2	+27.4%	<0.0001
<i>daf-2(e1370); daf-16(mgDf50)</i>	Insulin-like pathway	<i>E. coli</i> (OP50)	107	11.2 $\pm$ 0.1	<i>B. subtilis</i> (PY79)	114	12.3 $\pm$ 0.4	+9.9%	<0.001
<i>glp-1(e2141)</i>	Germ Line Proliferation	<i>E. coli</i> (OP50)	189	24.5 $\pm$ 0.8	<i>B. subtilis</i> (PY79)	203	29.8 $\pm$ 0.5	+22.0%	<0.0001
<i>isp-1(qm150)</i>	Electron transport chain	<i>E. coli</i> (OP50)	102	24.6 $\pm$ 0.8	<i>B. subtilis</i> (PY79)	174	32.0 $\pm$ 2.4	+30.2%	<0.0001
<i>nuo-6(qm200)</i>	Electron transport chain	<i>E. coli</i> (OP50)	213	31.9 $\pm$ 0.5	<i>B. subtilis</i> (PY79)	312	38.7 $\pm$ 1.8	+21.3%	<0.0001
<i>eat-2(ad1116)</i>	Dietary restriction	<i>E. coli</i> (OP50)	355	24.0 $\pm$ 0.6	<i>B. subtilis</i> (PY79)	408	28.2 $\pm$ 0.9	+17.3%	<0.0001
<i>hif-1(ia4)</i>	Hypoxia-induced factor	<i>E. coli</i> (OP50)	211	18.2 $\pm$ 0.6	<i>B. subtilis</i> (PY79)	249	22.3 $\pm$ 1.4	+22.1%	<0.0001
<i>hsf-1(sy441)</i>	Heat-shock transcription factor	<i>E. coli</i> (OP50)	107	13.1 $\pm$ 0.3	<i>B. subtilis</i> (PY79)	112	15.1 $\pm$ 0.4	+15.6%	<0.0005
<i>jnk-1(gk7)</i>	Jun N-terminal Kinase	<i>E. coli</i> (OP50)	118	15.0 $\pm$ 0.3	<i>B. subtilis</i> (PY79)	102	21.6 $\pm$ 0.6	+44.5%	<0.0001
<i>dbl-1(nk3)</i>	Transforming growth factor beta pathway	<i>E. coli</i> (OP50)	93	14.7 $\pm$ 0.4	<i>B. subtilis</i> (PY79)	94	21.0 $\pm$ 0.8	+42.5%	<0.0001
<i>pmk-1(km25)</i>	Mitogen-activated protein kinase	<i>E. coli</i> (OP50)	118	14.5 $\pm$ 0.5	<i>B. subtilis</i> (PY79)	128	19.1 $\pm$ 0.1	+32.0%	<0.0001
<i>tol-1(nr2033)</i>	Toll-like receptor	<i>E. coli</i> (OP50)	227	14.9 $\pm$ 0.3	<i>B. subtilis</i> (PY79)	222	29.7 $\pm$ 1.2	+98.9%	<0.0001



**Table S7. Calculations for proportional life span differences of mutants with respect to controls**

<b>*<i>C. elegans</i> strain</b>	<b>Median life span ± SD on <i>B. subtilis</i> diet</b>	<b>Number of life span assays (total number of worms)</b>	<b>**Projected median life span ± SD on <i>B.</i> <i>subtilis</i> diet</b>	<b>***p. value (T test) indicating probability that median life span is not proportionally similar to median life span of N2 controls</b>
Wild type (N2)	23.2±1.7	13 (465)		
<i>daf-16(mu86)</i>	15.7±1.3	6 (229)	18.0±1.5	p<0.0001
<i>daf-16(mgDf50)</i>	14.9±0.3	3 (115)	19.2±0.3	p<0.005
<i>daf-2(e1370); daf-16(mgDf50)</i>	12.3±0.8	3 (114)	17.0±1.0	p<0.0001
<i>glp-1(e2141)</i>	29.8±1.1	4 (203)	21.2±0.8	p<0.05
<i>eat-2(ad1116)</i>	24.3±1.6	9 (408)	20.0±1.3	p<0.0005
<i>hif-1(ia4)</i>	22.3±2.2	6 (249)	19.8±1.9	p<0.005
<i>hsf-1(sy441)</i>	15.1±0.8	3 (112)	18.0±0.9	p<0.0001
<i>pmk-1(km25)</i>	19.1±0.2	3 (128)	20.4±0.2	p<0.05
<i>tol-1(nr2033)</i>	29.7±3.0	6 (222)	30.8±3.1	p<0.0001

\* Strains used in Figure 6, which are not represented in this table, displayed median life spans that were proportionally similar to the median life span of N2 controls (p>0.05, T test)

\*\* Life span data for *B. subtilis* fed mutant worms after normalizing with respect to median life span reference point obtained for N2 worms fed *E. coli* (16.0 days) and *B. subtilis* (23.2 days)

\*\*\* T test compares normalized median life spans of *B. subtilis* fed mutant worms with median life spans of *B. subtilis* fed N2 wild type worms