

## SUPPLEMENTARY TABLES

**Supplementary Table 1. Statistical analyses of lifespan assays.**

| Genotype                                                                 | RNAi                  | Tissue <sup>a</sup> | Lifespan (days) |        | Percent of the control <sup>b</sup> | n <sup>c</sup> | p <sup>d</sup>   |
|--------------------------------------------------------------------------|-----------------------|---------------------|-----------------|--------|-------------------------------------|----------------|------------------|
|                                                                          |                       |                     | Mean            | Max    |                                     |                |                  |
| Effect of the <i>par-1(zu310)</i> mutant on lifespan (Figure 1A)         |                       |                     |                 |        |                                     |                |                  |
| N2                                                                       | /                     | /                   | 13.66, 11.71    | 19, 17 | /                                   | 79, 56         | /                |
| <i>par-1(zu310)</i>                                                      | /                     | /                   | 16.39, 14.53    | 25, 19 | 120%, 124%                          | 49, 68         | <0.0001, <0.0001 |
| Temporal requirement of <i>par-1</i> in lifespan regulation (Figure 1B)  |                       |                     |                 |        |                                     |                |                  |
|                                                                          | control               |                     | 13.93, 13.14    | 19, 17 | /                                   | 84, 58         | /                |
| N2                                                                       | <i>par-1</i> (Dev)    | global              | 14.23, 12.88    | 19, 17 | 102%, 98%                           | 60, 50         | 0.5368, 0.3680   |
|                                                                          | <i>par-1</i> (AD)     |                     | 17.18, 14.69    | 23, 19 | 123%, 112%                          | 80, 71         | <0.0001, <0.0001 |
|                                                                          | <i>par-1</i> (Dev+AD) |                     | 17.73, 14.74    | 25, 21 | 127%, 112%                          | 74, 69         | <0.0001, <0.0001 |
| Spatio requirement of <i>par-1</i> in lifespan regulation (Figure 3)     |                       |                     |                 |        |                                     |                |                  |
| N2                                                                       | control               | global              | 14.03, 13.63    | 19, 17 | /                                   | 70, 63         | /                |
|                                                                          | <i>par-1</i>          |                     | 16.28, 15.79    | 23, 21 | 116%, 116%                          | 69, 66         | <0.0001, <0.0001 |
| <i>rrf-1</i>                                                             | control               | germline +          | 13.32, 13.51    | 17, 17 | /                                   | 63, 55         | /                |
|                                                                          | <i>par-1</i>          | intestine           | 13.94, 13.76    | 19, 17 | 105%, 102%                          | 68, 58         | 0.1756, 0.2838   |
| <i>rde-1; kbIs7</i>                                                      | control               | intestine           | 12.36, 12.05    | 17, 15 | /                                   | 50, 59         | /                |
|                                                                          | <i>par-1</i>          |                     | 12.15, 11.97    | 17, 15 | 98%, 99%                            | 54, 60         | 0.6433, 0.6665   |
| <i>rde-1; nels9</i>                                                      | control               | muscle              | 13.67, 13.16    | 19, 17 | /                                   | 63, 61         | /                |
|                                                                          | <i>par-1</i>          |                     | 13.67, 13.44    | 19, 17 | 100%, 102%                          | 60, 64         | 0.7955, 0.6097   |
| <i>rde-1; kzIs9</i>                                                      | control               | epidermis           | 13.63, 13.44    | 19, 17 | /                                   | 63, 63         | /                |
|                                                                          | <i>par-1</i>          |                     | 15.43, 16.21    | 23, 21 | 113%, 121%                          | 69, 63         | <0.0001, <0.0001 |
| Epistatic analysis of <i>par-1</i> for its effect on lifespan (Figure 4) |                       |                     |                 |        |                                     |                |                  |
| <i>daf-16</i>                                                            | control               | global              | 9.61, 11.40     | 13, 17 | /                                   | 62, 177        | /                |
|                                                                          | <i>par-1</i>          |                     | 11.24, 13.06    | 17, 21 | 117%, 115%                          | 67, 174        | <0.0001, <0.0001 |
| <i>daf-2</i>                                                             | control               | global              | 28.82, 29.99    | 49, 39 | /                                   | 101, 75        | /                |
|                                                                          | <i>par-1</i>          |                     | 33.46, 33.64    | 53, 41 | 116%, 112%                          | 109, 84        | <0.0001, <0.0001 |
| <i>glp-1</i>                                                             | control               | global              | 15.24, 17.14    | 27, 23 | /                                   | 139, 58        | /                |
|                                                                          | <i>par-1</i>          |                     | 20.30, 18.97    | 27, 25 | 133%, 111%                          | 123, 61        | <0.0001, 0.0089  |
| <i>eat-2</i>                                                             | control               | global              | 16.77, 15.75    | 25, 21 | /                                   | 192, 69        | /                |
|                                                                          | <i>par-1</i>          |                     | 17.20, 17.36    | 25, 21 | 103%, 110%                          | 131, 67        | 0.1489, 0.0053   |
| <i>rsks-1</i>                                                            | control               | global              | 15.53, 14.94    | 21, 21 | /                                   | 60, 62         | /                |
|                                                                          | <i>par-1</i>          |                     | 15.39, 14.78    | 21, 21 | 99%, 99%                            | 61, 64         | 0.7661, 0.7359   |
| <i>aak-2</i>                                                             | control               | global              | 12.05, 12.30    | 15, 17 | /                                   | 61, 60         | /                |
|                                                                          | <i>par-1</i>          |                     | 11.87, 12.51    | 15, 17 | 99%, 102%                           | 60, 61         | 0.4647, 0.6463   |

<sup>a</sup>, tissues in which RNAi is mainly effective.

<sup>b</sup>, changes in mean lifespan compared to the control.

<sup>c</sup>, numbers of animals scored.

<sup>d</sup>, log-rank tests.

**Supplementary Table 2. Statistical analyses of thermotolerance assays.**

| Genotype | RNAi         | Survival (hours) |        | Percent of the control <sup>a</sup> | n <sup>b</sup> | p <sup>c</sup>   |
|----------|--------------|------------------|--------|-------------------------------------|----------------|------------------|
|          |              | Mean             | Max    |                                     |                |                  |
| N2       | control      | 15.31, 12.81     | 20, 18 | /                                   | 61, 47         | /                |
|          | <i>par-1</i> | 18.11, 15.70     | 22, 20 | 118%, 123%                          | 57, 54         | <0.0001, <0.0001 |

<sup>a</sup>, changes in the mean survival compared to the control.

<sup>b</sup>, numbers of animals scored.

<sup>c</sup>, log-rank tests.

**Supplementary Table 3. Statistical analyses of UV stress assays.**

| Genotype | RNAi         | Survival (days) |      | Percent of the control <sup>a</sup> | n <sup>b</sup> | p <sup>c</sup>  |
|----------|--------------|-----------------|------|-------------------------------------|----------------|-----------------|
|          |              | Mean            | Max  |                                     |                |                 |
| N2       | control      | 2.63, 2.90      | 4, 4 | /                                   | 54, 154        | /               |
|          | <i>par-1</i> | 3.24, 3.39      | 5, 6 | 123%, 117%                          | 59, 160        | 0.0003, <0.0001 |

<sup>a</sup>, changes in the mean survival compared to the control.

<sup>b</sup>, numbers of animals scored.

<sup>c</sup>, log-rank tests.

**Supplementary Table 4. Statistical analyses of proteotoxicity assays.**

| Genotype                             | RNAi         | Paralysis (days) |        | Percent of the control <sup>a</sup> | n <sup>b</sup> | p <sup>c</sup>   |
|--------------------------------------|--------------|------------------|--------|-------------------------------------|----------------|------------------|
|                                      |              | Mean             | Max    |                                     |                |                  |
| polyQ-induced paralysis              |              |                  |        |                                     |                |                  |
| <i>rmIs132(unc-54p::Q35::YFP)</i>    | control      | 9.48, 10.82      | 13, 15 | /                                   | 42, 113        | /                |
|                                      | <i>par-1</i> | 11.52, 12.40     | 15, 17 | 122%, 115%                          | 42, 100        | <0.0001, <0.0001 |
| Aβ-induced paralysis                 |              |                  |        |                                     |                |                  |
| <i>dvIs2(unc-54p::Aβ1-42 + pRF4)</i> | control      | 8.93, 10.52      | 14, 13 | /                                   | 69, 63         | /                |
|                                      | <i>par-1</i> | 11.24, 11.92     | 16, 15 | 126%, 113%                          | 68, 61         | <0.0001, <0.0001 |

<sup>a</sup>, changes in the mean paralysis time compared to the control.

<sup>b</sup>, numbers of animals scored.

<sup>c</sup>, log-rank tests.

**Supplementary Table 5. Statistical analyses of muscular function assays.**

| Age    | RNAi         | Number of body bends / 30"             | Average | STD  | n <sup>a</sup> | p <sup>b</sup> |
|--------|--------------|----------------------------------------|---------|------|----------------|----------------|
| day 2  | control      | 39, 35, 38, 42, 40, 41, 42, 40, 42, 45 | 40.4    | 2.72 | 10             | /              |
|        | <i>par-1</i> | 54, 52, 60, 50, 54, 50, 51, 52, 54, 60 | 53.7    | 3.65 | 10             | <0.0001        |
| day 4  | control      | 39, 35, 32, 33, 35, 33, 34, 35, 40, 35 | 35.1    | 2.56 | 10             | /              |
|        | <i>par-1</i> | 44, 45, 51, 45, 43, 46, 42, 43, 45, 43 | 44.7    | 2.54 | 10             | <0.0001        |
| day 6  | control      | 31, 26, 25, 26, 27, 32, 28, 30, 24, 30 | 27.9    | 2.73 | 10             | /              |
|        | <i>par-1</i> | 39, 40, 45, 35, 41, 36, 39, 35, 36, 46 | 39.2    | 3.94 | 10             | <0.0001        |
| day 8  | control      | 12, 20, 19, 14, 10, 16, 13, 18, 19, 18 | 15.9    | 3.45 | 10             | /              |
|        | <i>par-1</i> | 26, 27, 25, 25, 30, 26, 33, 28, 38, 26 | 20.6    | 4.20 | 10             | <0.0001        |
| day 10 | control      | 5, 1, 11, 0, 8, 11, 11, 2, 13, 6       | 6.9     | 4.61 | 10             | /              |
|        | <i>par-1</i> | 22, 26, 17, 20, 23, 23, 22, 20, 14, 19 | 20.6    | 3.41 | 10             | 0.0029         |

<sup>a</sup>, numbers of animals scored.

<sup>b</sup>, two-way ANOVA with Sidak's multiple comparison tests.