

**Table S2 List of sequences and sources for transgene construction**

**A. Primer sequences for GFP transcriptional fusions**

Regulatory region	Forward primer (5' to 3')	Reverse primer (5' to 3')
<i>cgh-1</i>	<i>SphI</i> - tttcacagattctgggtgaatatgagg	<i>KpnI</i> - ttcccgatgtcgtagttaggtttgatttc
<i>cpb-3</i>	<i>SphI</i> - accggaggaagcttggtgaagcaatg	<i>KpnI</i> - ggtgaatgcctttatcaacggc
<i>dcr-1</i>	<i>SphI</i> - aggattaaaatcttgaacatattggccg	<i>KpnI</i> - tattggctgaaaacaggaaaaattgat
<i>ddx-17</i>	<i>attB1</i> - gtatttcgtgacgtcactgcg	<i>attB2</i> - tttccctgtctccattgtc
<i>larp-5</i>	<i>attB1</i> - tccactttccattacgttgcggc	<i>attB2</i> - ctccggaaaaggggacgtcgacgccc
<i>mbl-1</i>	<i>attB1</i> - tttgggattacttcggtgctc	<i>attB2</i> - ccgatcgtgcgcattgtcc
<i>mtr-4</i>	<i>SphI</i> - tcgccacgaaaccacaccaaacggcgag	<i>KpnI</i> - ttatgtctaaaaatgtgccacaataaaa
<i>rsp-3</i>	<i>attB1</i> - gccctatcttaaatggccgg	<i>attB2</i> - gtttagtacactgaaaatgaag
<i>rsp-6</i>	<i>attB1</i> - gtgtaaatgtgatgcattcgag	<i>attB2</i> - actgaaaaatcaaattaatttatg
<i>set-2</i>	<i>attB1</i> - tgaatagcaaacttcataatcc	<i>attB2</i> - ttcccatcaattatttattgatgc
<i>Y55F3AM.3</i>	<i>attB1</i> - tcaaattgtatgttctattcgc	<i>attB2</i> - ggagctgaccccgcttcaaa

For *sup-26*, an existing transgene, *smls259[P<sub>sup-26</sub>::sup-26 cDNA::GFP]*, was used (Mapes *et al.* 2010).

**B. Primer sequences for GFP translational fusions to GFP**

cDNA (source)	Forward primer (5' to 3')	Reverse primer (5' to 3')
<i>cgh-1</i> (yk85e1, yk1658a03)	<i>attB1</i> - atgagtggagcggagcaacaacag	<i>attB2</i> - agcagtggctcatcgccggc
<i>cpb-3</i> (this study)	<i>attB1</i> - atgaacctgaatgaccgagtggaaag	<i>attB2</i> - accggaggaagcttggtgaagcaatg
<i>ddx-17</i> (this study)	<i>attB1</i> - atggagacagaggatacggagg	<i>attB2</i> - ccatcgccacctccaccgctgc
<i>larp-5</i> (yk1291c05, yk400c5)	<i>attB1</i> - atggacacgttcgagctcgactcg	<i>attB2</i> - cctgaacgtcggttttgtggcgg
<i>mbl-1</i> (this study)	<i>attB1</i> - atgttcgacgaaaacagtaatgccg	<i>attB2</i> - gaatgggtggctgcatgtac
<i>mtr-4</i> (yk506b7, yk430a11)	<i>attB1</i> - atggccgatttcgacgaattc	<i>attB2</i> - tagatagagagatgcagcgaaaac
<i>rsp-3</i> (yk1100h10, yk1195c10)	<i>attB1</i> - atgccacgcggcggctcagagg	<i>attB2</i> - ttgtggagatggtagcgagatgg

<i>rsp-6</i> (this study)	<i>attB1</i> - atggacgccaagggtgtacgtcg	<i>attB2</i> - gtgcggagaagcagaacggctgc
<i>set-2</i> (yk21g8, yk320e10)	<i>attB1</i> - atgtccacacatgatgatgaacc	<i>attB2</i> - attaagatatccacgacacgtc
<i>sup-26</i> (D. Xue, J. Mapes)	<i>attB1</i> - atgaacgcacatctcggtccac	<i>attB2</i> - ttgtggattcatcggtggaaatac
<i>Y55F3AM.3</i> (this study)	<i>attB1</i> - atgaccgacggatcgatgg	<i>attB2</i> - atatctgccatatccaccatattg

### C. Primer sequences for RNAi vectors

RNAi target	Forward primer (5' to 3')	Reverse primer (5' to 3')
<i>B0511.6</i>	<i>attB1</i> - gcattgacgtcaagaagaagggtgc	<i>attB2</i> - ccaccatactaatccatacg
<i>C44B7.2</i>	<i>attB1</i> - atgcaaggaaaggaggcgatcatcacg	<i>attB2</i> - atctggaagagtgaagtctcggtgatc
<i>cyn-13</i>	<i>attB1</i> - aatttccagctccgccacacaatgcccc	<i>attB2</i> - taagtctccatagcctgcggaaagcggc
<i>ddx-17</i>	<i>attB1</i> - tggctgcccggagaaccta	<i>attB2</i> - caattcacgagtttggaaagca
<i>rps-3</i>	<i>attB1</i> - cgtgaccaagaagaagaaggccg	<i>attB2</i> - gatgttagtcgttgactgggtgtcc
<i>Y23H5B.6</i>	<i>attB1</i> - tatgcctcacacaaacggaaaaggcggcg	<i>attB2</i> - atttgtcaaactctgcacaaatccctg
<i>ZC190.4</i>	<i>attB1</i> - tatcttgatactggctgaaaaagttgcg	<i>attB2</i> - tacacaactgggtatgtgaagggcac
<i>ZC434.3</i>	<i>attB1</i> - tatgcgtctcgtaaaaccactgtatgtgg	<i>attB2</i> - ctagttttctgagttgtctcgaaaatc

Genomic fragments PCR-amplified with primers sequences given were cloned into pDONR221 with a BP reaction and then cloned into a double-T7 plasmid with a Gateway entry site (modified from pPD129.36) using an LR reaction (Invitrogen).