

Table S1. Primers used in this study

	Forward	Reverse
Cloning		
<i>daf-8::gfp</i>	GATAGGTAGGAAAAAGCAACGAAG	GAAAAGTTCTTCTCCTTTACTCATAGTTCTGGATGAACATATACG
<i>daf-8::FLAG</i>	GATAGGTAGGAAAAAGCAACGAAG	TTGGATCCTCAGATCTTATCGTCGTCATCCTTGTAATCAGTTCTGGATGAACATATACG
<i>daf-14::FLAG</i>	GGGCTCGAGCTGTCTGACTTTTCTGGACTGCTCTAC	TTAAGCTTTCAGATCTTATCGTCGTCATCCTTGTAATCTGATCGGGATCCTATTTCCGG
<i>daf-3::FLAG</i>	AAACTCGAGCCGAGGTCATGTGCACTCTCTTG	TTTCTCGAGTCAGATCTTATCGTCGTCATCCTTGTAATCGTCATCACTGACGTTGATTCC
<i>daf-8 cDNA in pBSIIKS</i>	CCCGGATCCATGGACGATTTTCCTTCACCAG	TTTGAATTCCTAAGTTCTGGATGAACATATAC
<i>daf-8 cDNA in pGEX4T-1</i>	CCCGGATCCGACGATTTTCCTTCACCAGAACC	TTTGAATTCCTAAGTTCTGGATGAACATATAC
<i>daf-14 cDNA in pBSIIKS</i>	AAACTCGAGATGTACCCAGATCAAGTTCAC	AAACTCGAGTCATGATCGGGATCCTATTTCCG
<i>daf-14 cDNA in pGEX4T-1</i>	AAACTCGAGTACCCAGATCAAGTTCACTTAC	AAACTCGAGTCATGATCGGGATCCTATTTCCG
<i>daf-3 cDNA in pGEX4T-1</i>	TTTCTCGAGAAGCTAATAGCAACTTCTCTTC	TTTCTCGAGTTAGTCATCACTGACGTTG
qPCR		
<i>daf-7</i>	AACGAAATTCTCGACCAGCTGA	TGGATCTTTGGCGGTGTAGAAT
<i>daf-1</i>	CAGCAGTCATCCACGTGGCT	TCCGTCTCGCGTAGAGGATCT
<i>daf-4</i>	GGAAAAGTTTTCAAGGCGCAA	CGCACACAAATTCGACGATACTT
<i>daf-8</i>	CGTCTACTGTCTCACGAATGTGC	GGAGCTCCCATACAAGTGAGTACAC
<i>daf-3</i>	CCGAACTTGTACGGGTTTCCAAC	TGCCTTGTTGGGACGTATGGTTT
<i>lag-2</i>	TTCCTCTTACTCCTCACCTGCCTCC	GCGAGTGGGACTCTGTGACAATTC
<i>act-2</i>	GGACTTGTACGCCAACACTG	CACATCTGTTGGAAGGTGGA
ChIP		
<i>daf-7 BD2</i>	GCAAAACGGCCTAACTTCAT	CGAAATTTAGCTTTGCATCTGC
<i>daf-8 BD1</i>	CTTACTCCGTAATGTTGGCATTTC	TGGCTTCAAAGACTGGTCAGGT
<i>daf-8 BD3</i>	CTTTAAAAAAGAGCAGAGTTGAGAG	GGCTAATTTGAGAGACACATTCTAA