

## Supplementary Table 2

*Primer sequences used for C. elegans qRT-PCR*

Gene	Forward (5' to 3')	Reverse (3' to 5')
<i>mlc-2</i>	TTGACAGGAACTGACCCAGAGG (265 bp to 286 bp)*	ATAGCCTTGACCTCATCCTCG (413 bp to 393 bp)
<i>cdr-1</i>	TCTTCTCTCAATTGGCAACTG (698 bp to 718 bp)	TTTGGGTAACTTCATGACGA (818 bp to 798 bp)
<i>numr-1/-2**</i>	GGAAGACGAGGAATGGATCATC (682 bp to 703 bp)	CTGCGTTCAAATGGATGTTGTCTC (830 bp to 807 bp)
<i>rpb-12</i>	CGCCGAAAATGAAATCAAAC (81 bp to 100 bp)	GGGCGTCGTACACCATCA (184 bp to 167 bp)

\*Sequence location of the primers using ATG as nucleotide 1.

\*\* Sequences for the coding regions of *numr-1* and *numr-2* are 100% identical, therefore the *numr*-specific primers will amplify both *numr-1* and *numr-2* cDNAs. For this reason the genes are designated *numr-1/-2*.