

Table S1. Conservation of proximal promoters and distal sequence among four nematode species

Species	length (bp) of proximal promoter	base pairs of proximal sequence in blocks ≥ 10 nt 100% conserved with <i>C. elegans</i>	% proximal sequence in blocks ≥ 10 nt 100% conserved with <i>C. elegans</i>	length of distal sequence used in assays	base pairs of distal sequence in blocks ≥ 10 nt 100% conserved with <i>C. elegans</i>	% distal sequence in blocks ≥ 10 nt 100% conserved with <i>C. elegans</i>
<i>C. briggsae</i>	254	82	32.3%	1121	11	1.0%
<i>C. brenneri</i>	242	92	38.0%	730	0	0.0%
<i>C. remanei</i>	230	88	38.3%	923	11	1.2%