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

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