

**Dataset S2.**

*Caenorhabditis elegans* N2 insertion-deletion (indel) summary statistics after an average of 250 generations (Gen). SEM is the standard error.

MA Line	Indels		Sites ( $\times 10^6$ )	Gen.	Indel Rate ( $\times 10^{-10}$ ) /events /gen.
	Ins.	Del.			
Ce 523	5	8	64.5	247	8.16
Ce 526	3	0	60.5	245	2.03
Ce 529	9	4	82.4	256	6.16
Ce 538	9	8	83.5	250	8.16
Ce 535	0	0	56.2	256	0.00
Ce 553	3	3	77.5	241	2.68
Ce 574	21	8	86.5	258	13.0
Sum	50	31	--	--	--
Ave.	7.14	4.43	73.01	250	5.74
SEM	2.83	1.49	5.03	2.63	1.83