

Table S2. Initial expansion size of trinucleotide repeat tracts^a

Genotype	Tract sequence	Median size (net change)^b	Range (net change)^b
<i>MSH3</i> (n=61)	<i>CAG</i>	42 (+17)	31-67 (+6 to +42)
<i>MSH3</i> (n=89)	<i>CTG</i>	33 (+8)	26-43 (+1 to +18)
<i>msh3Δ</i> (n=38)	<i>CAG</i>	31 (+6)	31-48 (+6 to +16)
<i>msh3Δ</i> (n=35)	<i>CTG</i>	33.5 (+8)	28-39 (+3 to +14)

^a Initial expansion sizes obtained from all time course experiments.

^b Number of repeats in TNR tract