



Supplemental Figure S13.

Strand asymmetry of deletions. Vertical axis, the ratio of the single nucleotide deletions rate in polyA tracts to single nucleotide deletions rate in polyT tracts in 10 MSI genomes (3 COAD samples and 7 UCEC samples). Notations as in Fig. 1. **a**, tract length 6 bp; **b**, tract length 7 bp; **c**, tract length 8 bp.