

Supplemental Table S7.

Replication asymmetry in MSI cancers in the most extreme bin and in the entire dataset (data for genomes samples from TCGA database). The values in brackets are 95% confidence intervals.

Asterisks denote significance of the deviation from 1.

*p:<0.05, **:p< 0.001,***:p<0.001.

Mutation	Rightmost bin	Entire dataset
A→G/T→C	1.83*** (1.79-1.87)	1.28*** (1.27-1.29)
A→C/T→G	1.50*** (1.44-1.58)	1.18*** (1.15-1.20)
A→T/T→A	0.92*** (0.88-0.96)	0.97** (0.95-0.99)
C→G/G→C	1.04 (0.96-1.12)	1.01 (0.97-1.04)
DpC→DpA/GpH→TpH	0.83*** (0.79-0.87)	0.92*** (0.90-0.94)
CpC→CpA/GpG→TpG	0.48*** (0.46-0.51)	0.74*** (0.73-0.76)
HpC→HpT/GpD→ApD	1.15*** (1.13-1.18)	1.06*** (1.04-1.07)
GpC→GpT/GpC→ApC	1.50*** (1.45-1.54)	1.18*** (1.16-1.19)