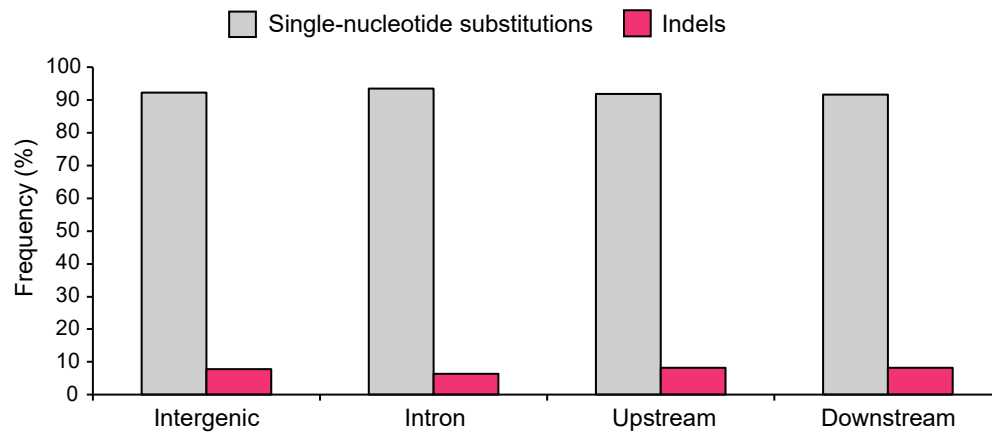




Supplementary Figure S1. Relative frequency of single-nucleotide substitutions (SNS) and indels in various non-coding regions of the genome of 2809 tumors analyzed by whole genome sequencing (PCAWG study available at <https://xenabrowser.net/datapages/>)