



S1 Figure. TC dinucleotide sequences are significantly depleted in PV genomes. The O/E ratios of all dinucleotides in PV genome sequences were analyzed with genome sequences of HPVs (A, $n = 161$), non-human primate (NHP) PVs (B, $n = 16$), and non-primate (NP) PVs (C, $n = 97$). Box-and-whisker plots are shown with the outliers (black triangles) identified using Tukey's method.