S8 Fig. Endogenous reverse transcription data. (A) Representative viral mutation plots from 1 of 4 independent ERT experiments. The *pol* region was amplified by high-fidelity PCR, cloned, and sequenced. (B) Actual distribution of G-to-A mutations in the indicated dinucleotide contexts in 10 independent 564 bp *pol* region DNA sequences from panel A.