

Figure S1. The relative and cumulative inertia of the first 40 axes accounting for 100% of the 59 axes.

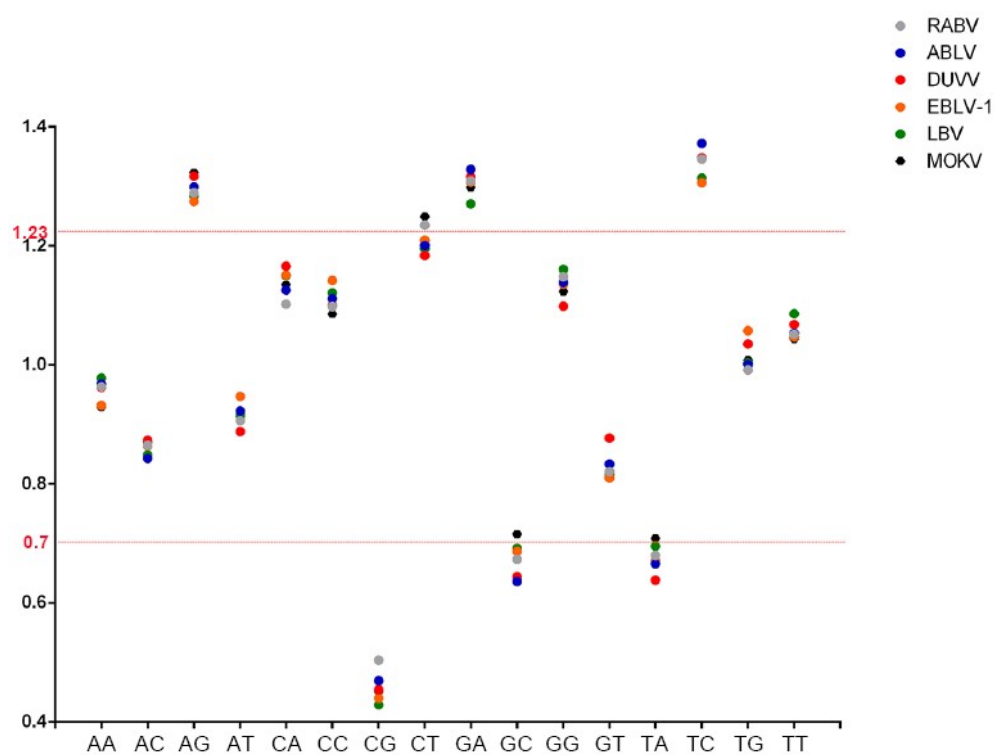


Figure S2. Dinucleotide frequency analysis. Table S1. Characteristic of lyssavirus genomes. Table S2. Nucleotide composition analysis.