



Figure 5 Predicted barbell-like structure of partial 3'UTR of MuKV/XM34/CHN and sequence alignment of partial 3'UTRs of murine kobuvirus strains. The sequence alignment in the 49-nt-long partial 3'UTRs begins after the stop codon, and the numbers on both sides of the sequences indicate the nucleotide position. The reference *Kobuvirus* strains include Wencheng-Rt386-2 (accession no.MF352432.1), rat08/rAiA/HUN (accession no.MN116647.1), MKV1/NYC/2014/M014/0146 (accession no.MF175074.1), M-5/USA/2010 (accession no.JF755427.1) and TF5WM (accession no.JQ408726.1).