Figure 1  Phylogenetic relationships of the three isolates and other strains including reference ALV-Js and other retroviruses based on U3 5′LTR (Figure 1A), 5′UTR (Figure 1B) and env gene (Figure 1C). Isolates marked with ▲ are three isolates in our study, and strains marked with ■ are also hemangioma ALV-Js from GenBank.