**Supplementary data 3 A-L:** Coding nucleotide (nt) and deduced amino acid (aa) sequence identities (%) for the A. VP1, B. VP2, C. VP3, D. VP4, E. VP6, F. VP7 (G2), G. VP7 (G8), H. NSP1, I. NSP2, J. NSP3, K. NSP4 and L. NSP5 genes of the study strains compared to those of human and animal strains from GenBank.

**Supplementary data 3 A.** VP7 (G8)



**Supplementary data 3 B.** VP7 (G2)



**Supplementary data 3 C.** VP4



**Supplementary data 3 D.** VP1



**Supplementary data 3 E.** VP2



**Supplementary data 3 F.** VP3



**Supplementary data 3 G.** VP6



**Supplementary data 3 H.** NSP1



**Supplementary data 3 I.** NSP2



**Supplementary data 3 J.** NSP3



**Supplementary data 3 K.** NSP4



**Supplementary data 3 L.** NSP5

