

Erratum to: Hepatitis E Virus Genotype 4 Sequences Detected in Sewage from Treatment Plants of China

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The original version of this article unfortunately contained a mistake in Fig. 1. Position of G3 and G4 was reversed. The corrected Fig. 1 is given below.

The online version of the original article can be found under
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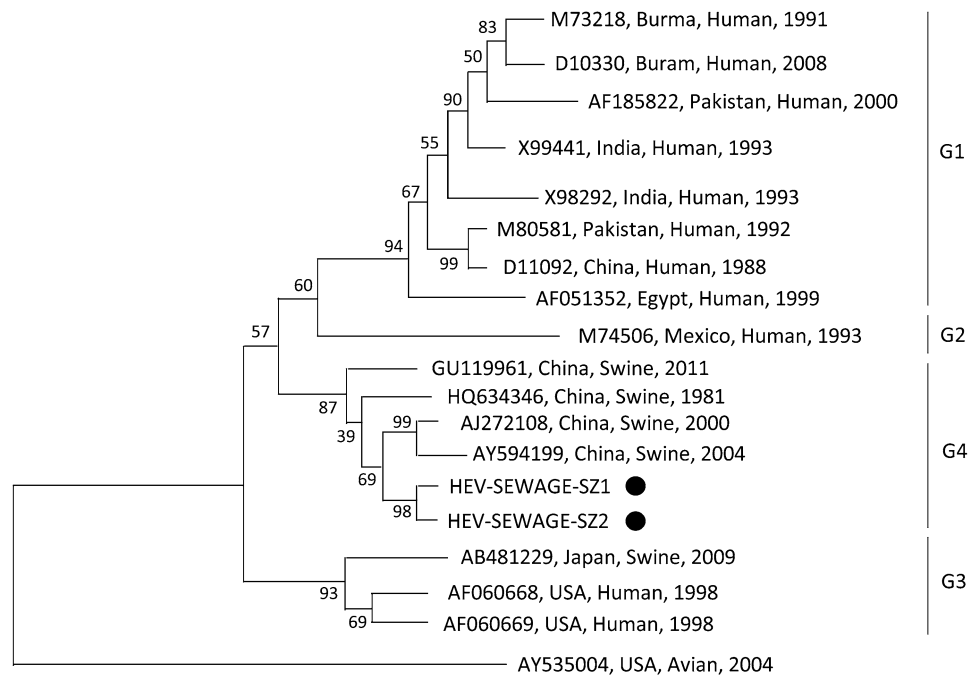
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Fig. 1 Phylogenetic analysis based on ORF2 (348 nt) depicting the genetic relationship between our strain from this study. A neighbor-joining tree was constructed with bootstrap values calculated from 1000 replicates. Isolates used for comparative analysis were Burma (GenBank Accession No. M73218, D10330), Pakistan (AF185822), India (X99441), India (X98292), Pakistan (M80581), China (D11092), Egypt (AF051352), Mexico (M74506), China (GU119961), China (HQ634346), China (AJ272108), China (AY594199), Japan (AB481229), USA (AF060669), USA (AF060668), and Avian HEV (AY535004)



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