Hepatitis C Virus Infection Caused by Infrequent Exposure in China Should Be of Concern

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Supplementary fig. 1  Details of the phylogenetic relationships of the partial NS5 gene. Bootstrap values >50% are shown at relevant nodes. Branches are colored according to their geographic origins, indicated at the right side. Symbol ● indicates the two sisters’ sequences, ▲ indicates the top BLAST hits for the elder sister, ■ indicates the top BLAST hits for the younger sister. The tree is midpoint rooted for clarity. The sequences of NS5b gene obtained here have been submitted to GenBank and assigned the accession numbers MH463463–MH463464.
Supplementary fig. 2  Details of the phylogenetic relationships of the partial E1 gene. Bootstrap values >50% are shown at relevant nodes. Branches are colored according to their geographic origins, indicated at the right side. Symbol ● indicates the two sisters’ sequences, ▲ indicates the top BLAST hits for the elder sister, ■ indicates the top BLAST hits for the younger sister. The tree is mid-point rooted for clarity. The sequences of E1 gene obtained here have been submitted to GenBank and assigned the accession numbers MH463465–MH463466.