

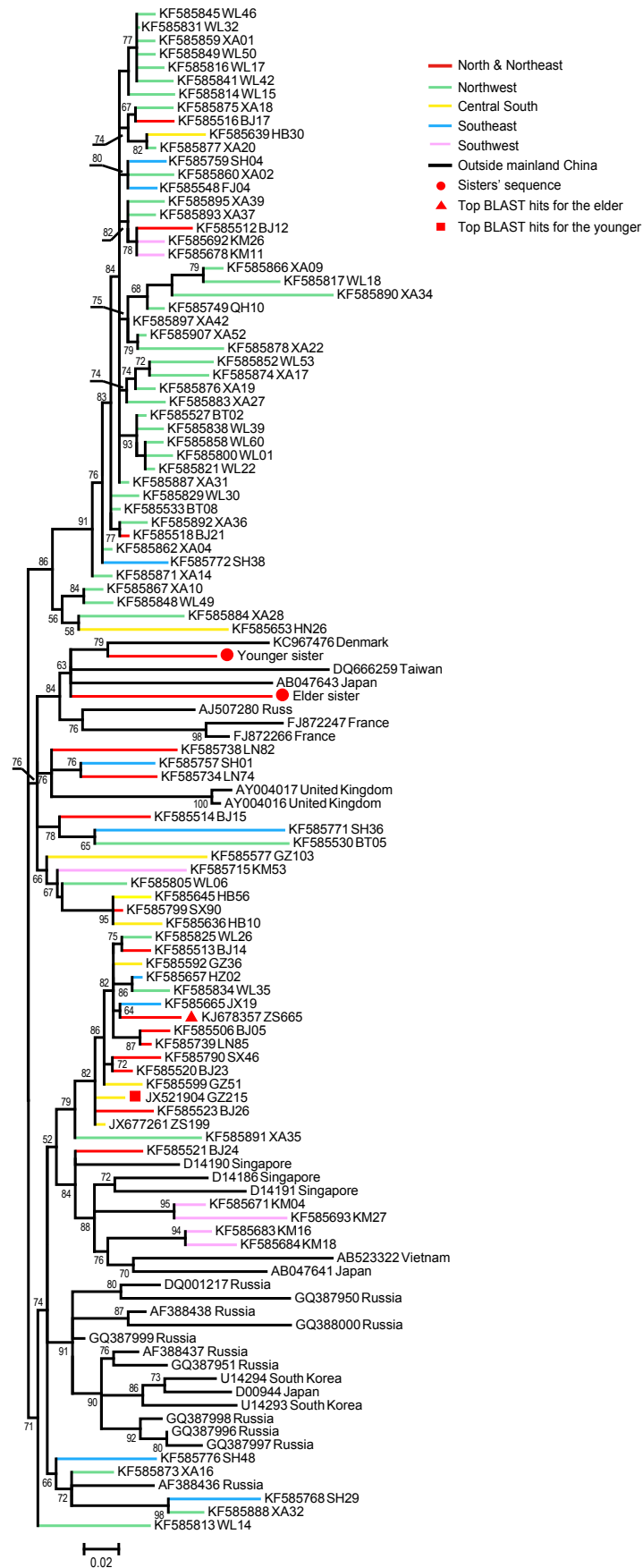
## Electronic Supplementary Material

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

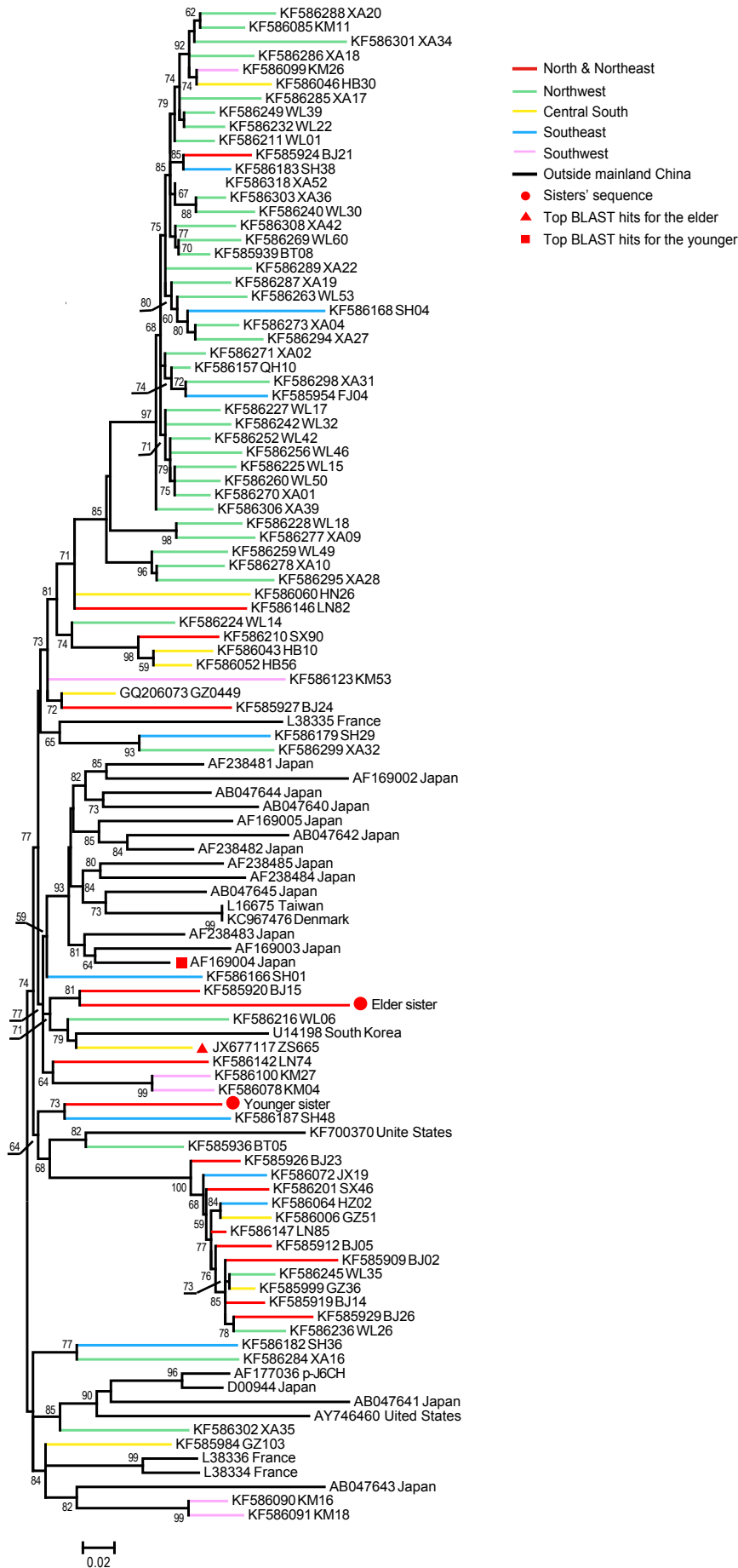
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Supplementary fig. 1 Details of the phylogenetic relationships of the partial *NS5* gene. Bootstrap values >50% are shown at relevant nodes. Branches are colored according to their geographic origins, indicated at the right side. Symbol ● indicates 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 mid-point 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 *EI* 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 *EI* gene obtained here have been submitted to GenBank and assigned the accession numbers MH463465–MH463466.