

## Electronic Supplementary Material

### **Spatial, Temporal and Genetic Dynamics Characteristics of Influenza B Viruses in China, 1973–2018**

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Table S1. Accession numbers of HA gene sequences used in study for Victoria lineage viruses phylodynamics in China.

Table S2. Accession numbers of HA gene sequences used in study for Yamagata lineage viruses phylodynamics in China.

Table S3. Summary of outbreak and sequence data by year during 2008–2018.