**Supplementary Materials**

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**Individual virome analysis reveals the general co-infection of mammal-associated viruses with SARS-related coronaviruses in bats**

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**Supplementary Tables**

**Excel files**

**Supplementary Table S1.** Sample information and reference studies for SARSr-CoV positives.The statistical results corresponding to Fig. 1B are shown in the attached tables.

**Supplementary Table S2.** Mapping results of potential co-circulating viruse**s.**

**Supplementary Table S3.** Mapping results of potential co-circulating fungi.

**Supplementary Table S4.** Mapping results of potential co-circulating bacteria.

**Supplementary Table S5.** GenBase accession of the 25 new viruses.

**Supplementary Table S6.** Metadata of reference sequences.

**Supplementary Table S7.** Blast results of the 25 new viruses.

**Supplementary Table S8.** Genome identity of the 25 new viruses.