Epidemiologic Evidence for Airborne Transmission of SARS-CoV-2 during Church Singing, Australia, 2020

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Appendix Figure 2. Phylogenetic analysis of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) genomes sequenced from case-patients during outbreak in the church, Australia, 2020. Black node indicates the tree root (GISAID accession no. EPI-ISL-413856); red node indicates primary case-patient; blue node indicates secondary case-patient from July 15 service; and green node indicates secondary case-patient from July 16 service. Scale bar indicates nucleotide substitutions per site. GISAID virus names of SARS-CoV-2 genomes hCoV-19/Australia/NSW657/2020, hCoV-19/Australia/NSW682/2020, hCoV-19/Australia/NSW698/2020, hCoV-19/Australia/NSW699/2020, hCoV-19/Australia/NSW707/2020, hCoV-19/Australia/NSW708/2020, hCoV-19/Australia/NSW715/2020, hCoV-19/Australia/NSW727/2020, hCoV-19/Australia/NSW747/2020, hCoV-19/Australia/NSW755/2020.
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