Recombination of Human Coxsackievirus B5 in Hand, Foot, and Mouth Disease Patients, China

Technical Appendix

![Graph showing similarity scanning analysis of coxsackievirus B (CBV) strain CBV5/CC10/10 with other CBV strains and representative enterovirus strains on the basis of full-length genome sequences.]

Technical Appendix Figure 1. Similarity scanning analysis of coxsackievirus B (CBV) strain CBV5/CC10/10 with other CBV strains and representative enterovirus strains on the basis of full-length genome sequences.
Technical Appendix Figure 2. Full-length bootscanning analysis of coxsackievirus B (CBV) strain CBV5/CC10/10 with CBV3-Beijing0811, CBV5–2000/CSF/KOR, and CBV5-COXB5/Henan/2010. Bootscanning was performed with a window size of 200 nt and step of 20 nt.