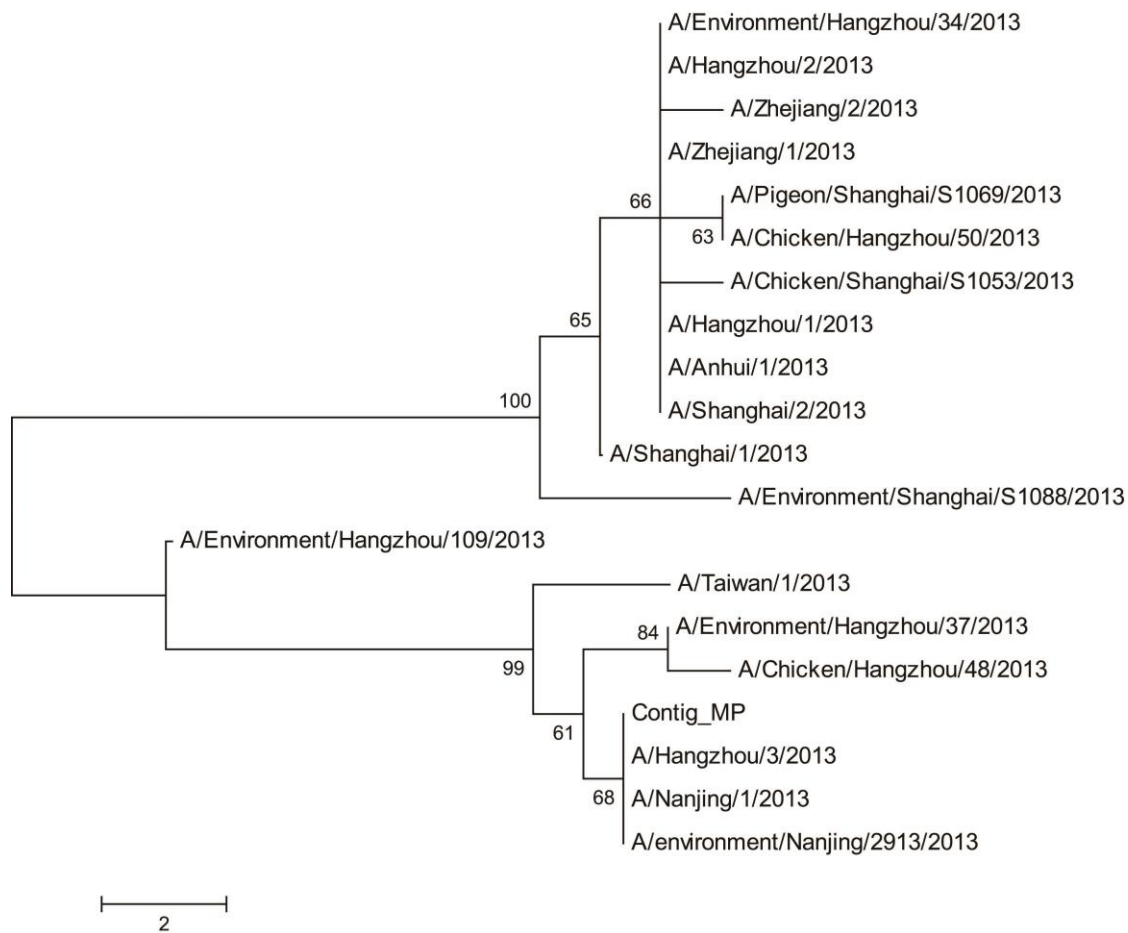
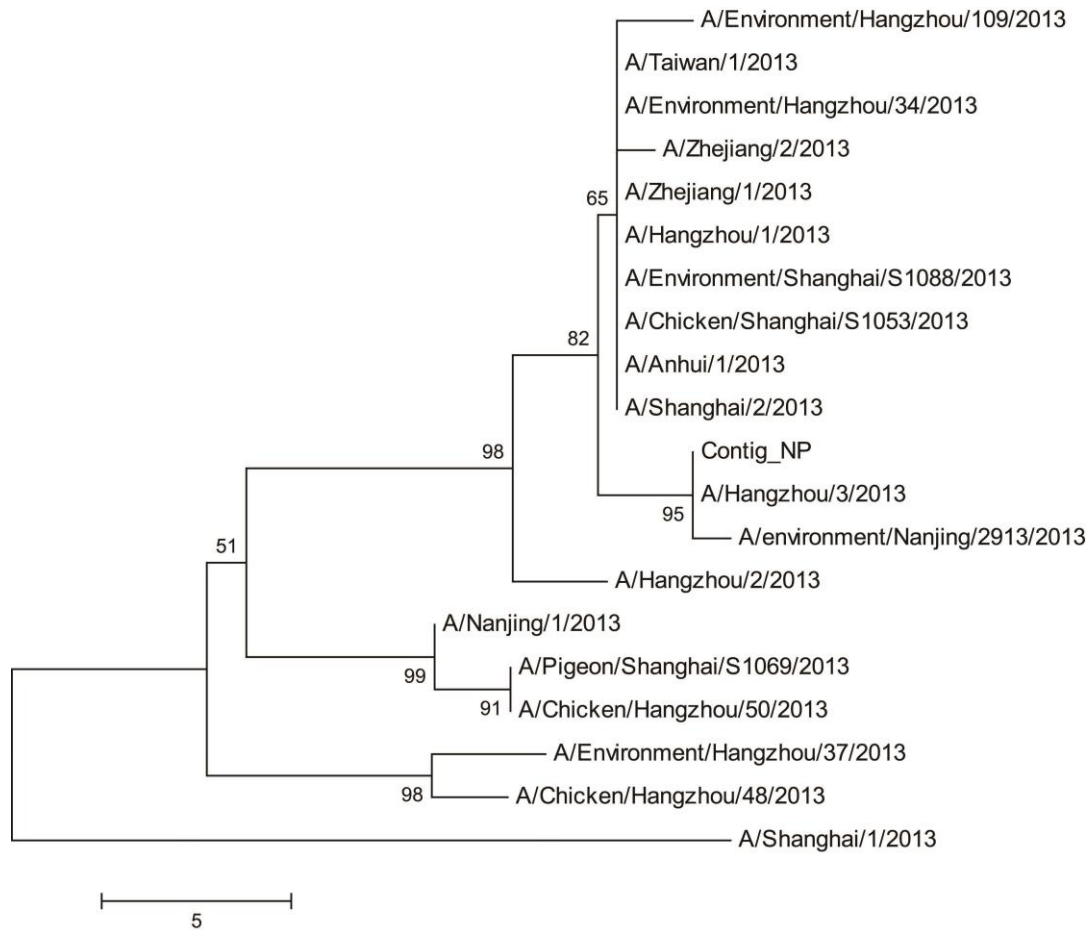


# Full Genome of Influenza A (H7N9) Virus Derived by Direct Sequencing without Culture

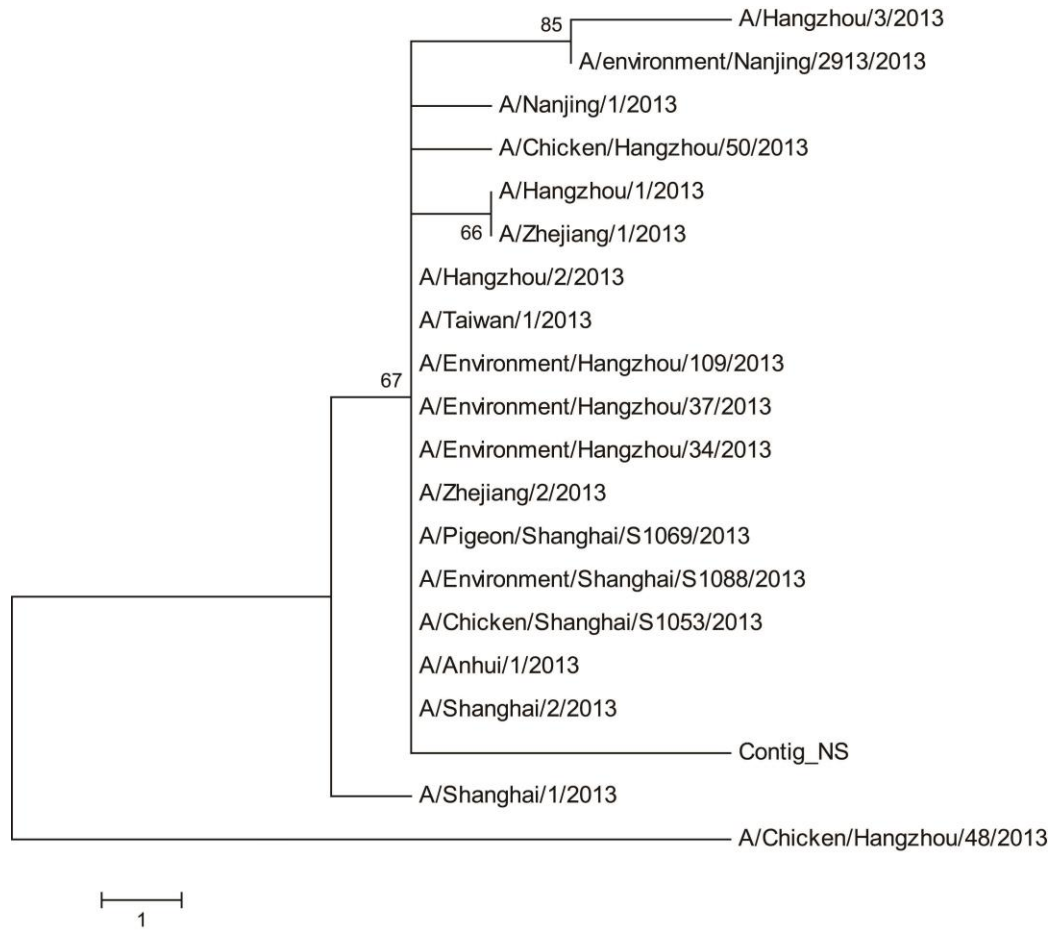
## Technical Appendix



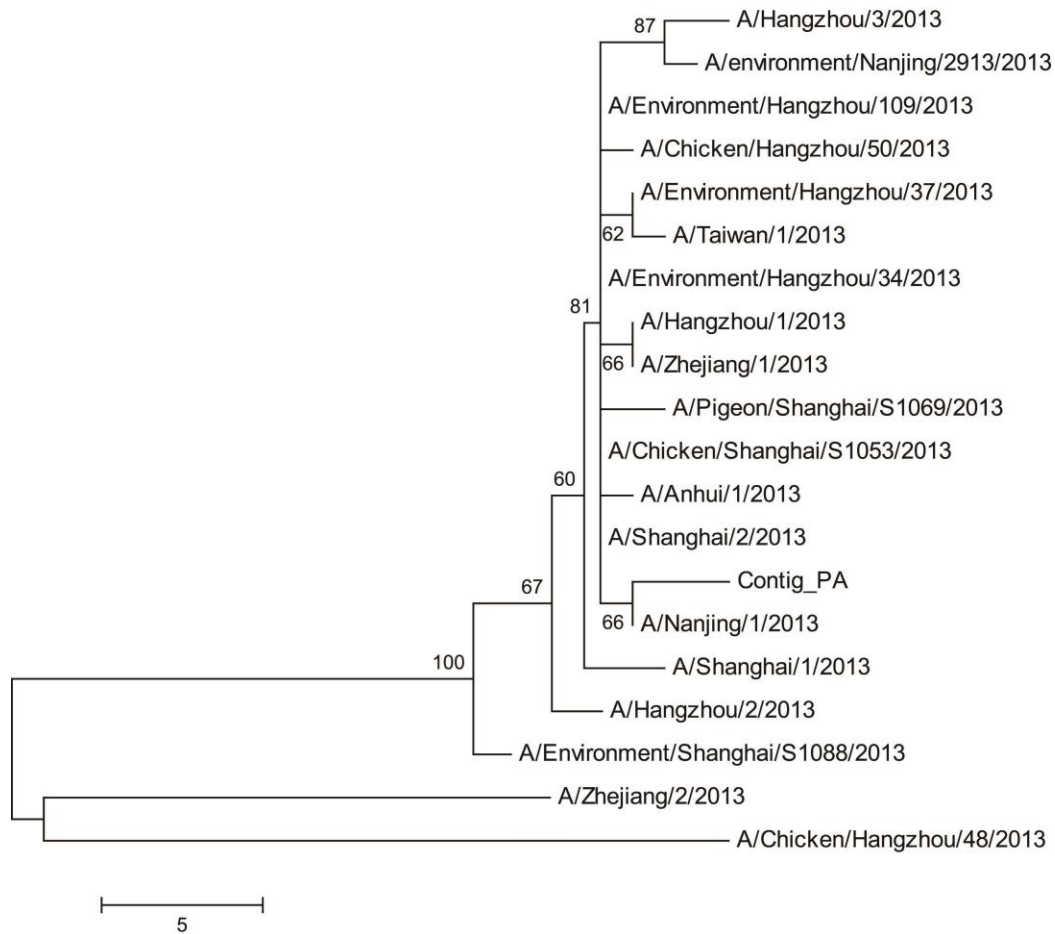
Technical Appendix Figure 1. Phylogenetic tree of the influenza A (H7N9) viruses isolated in China in 2013 based on the matrix protein gene segment. Scale bar indicates nucleotide differences per unit length.



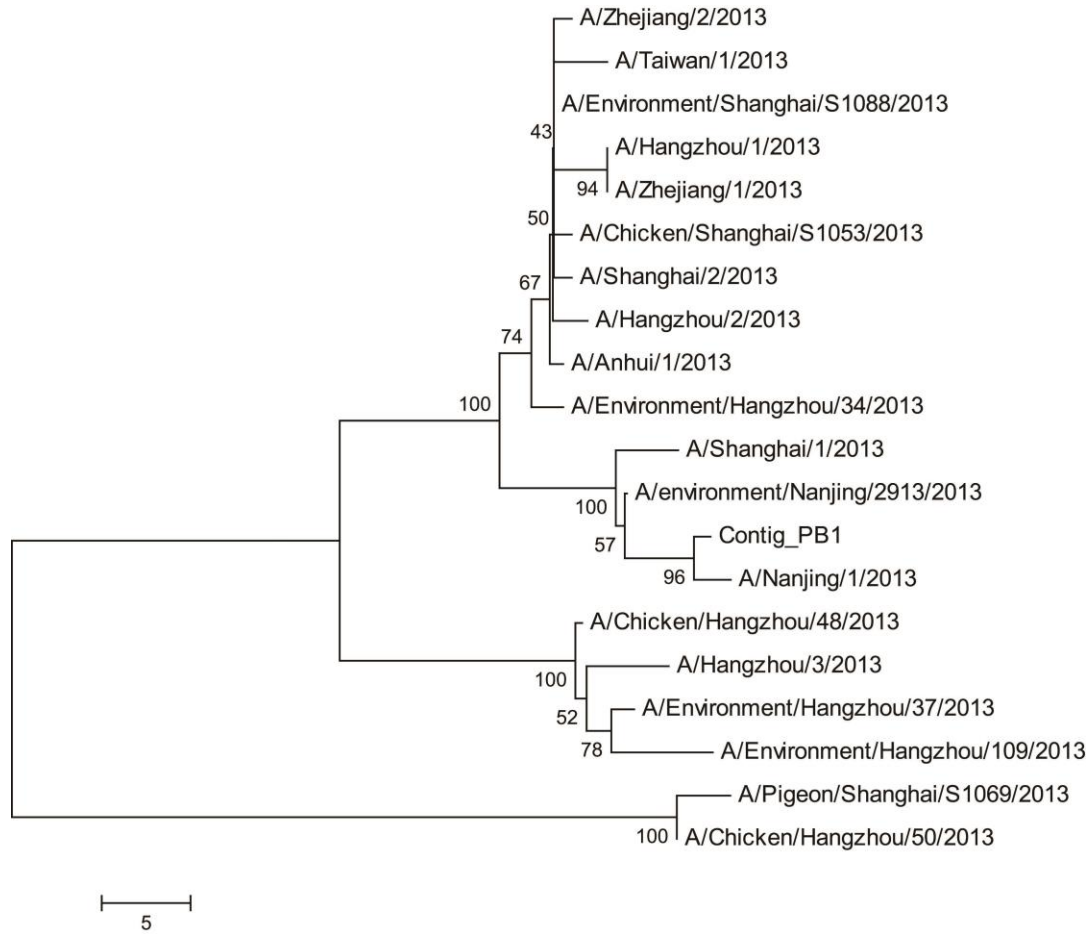
Technical Appendix Figure 2. Phylogenetic tree of the influenza A (H7N9) viruses isolated in China in 2013 based on the nucleocapsid protein gene segment. Scale bar indicates nucleotide differences per unit length.



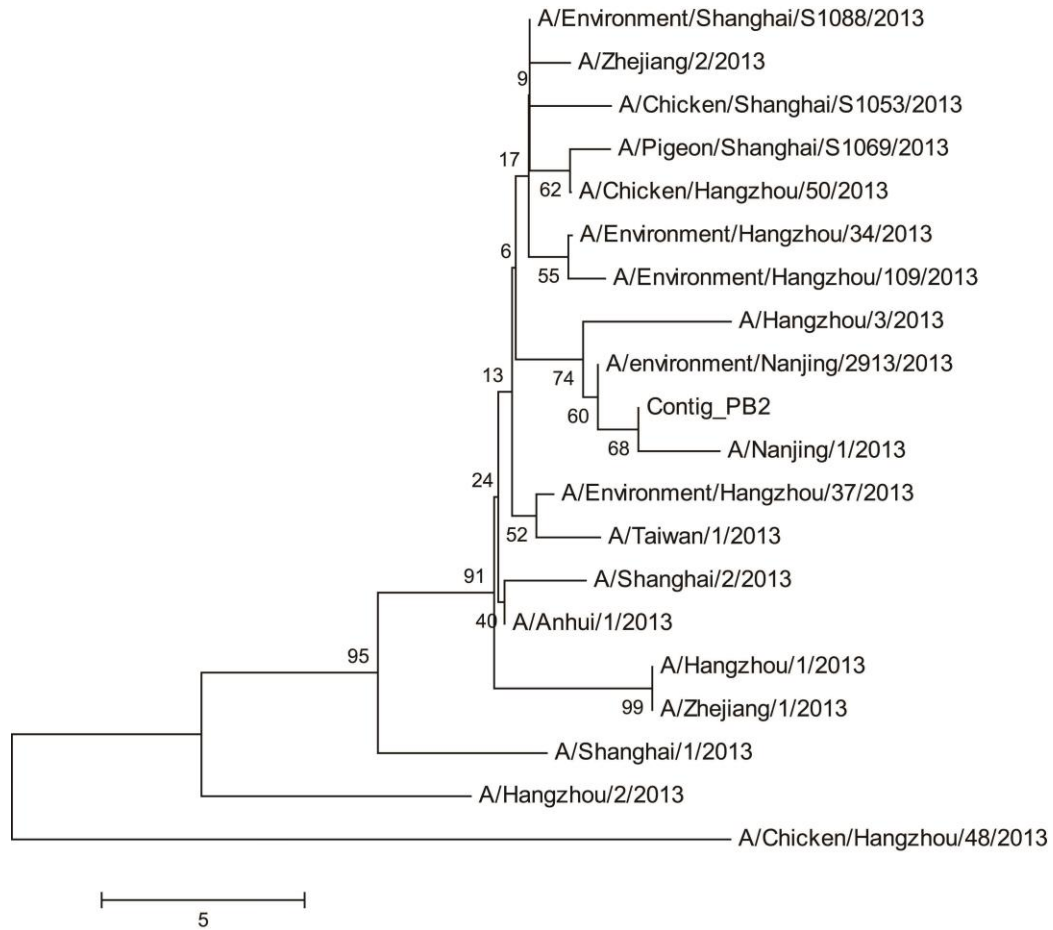
Technical Appendix Figure 3. Phylogenetic tree of the influenza A (H7N9) viruses isolated in China in 2013 based on the nonstructural protein gene segment. Scale bar indicates nucleotide differences per unit length.



Technical Appendix Figure 4. Phylogenetic tree of the influenza A (H7N9) viruses isolated in China in 2013 based on the polymerase acidic protein gene segment. Scale bar indicates nucleotide differences per unit length.



Technical Appendix Figure 5. Phylogenetic tree of influenza A (H7N9) viruses isolated in China in 2013 based on the polymerase basic protein 1 gene segment. Scale bar indicates nucleotide differences per unit length.



Technical Appendix Figure 6. Phylogenetic tree of the influenza A (H7N9) viruses isolated in China in 2013 based on the polymerase basic protein 2 gene segment. Scale bar indicates nucleotide differences per unit length.