Figure 2. Phylogenetic tree of partial RNA polymerase sequences of the viral strains detected in the food handler and five cases, and reference strains.

The tree was constructed using the UPGMA clustering method. Bootstrap values (1,000 replicates) are shown at the branch nodes. The food handler’s sample is labeled with a triangle and the five cases’ specimens are labeled with round circles.

The sequences of reference strains are from the Foodborne Viruses in Europe database (http://www.eufoodbornviruses.co.uk/).