

Corrigendum

Sequence of 3,687 nucleotides from the 3' end of Sendai virus genome RNA and the predicted amino acid sequences of viral NP,P and C proteins

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In Fig.2 on page 7,322, the sequence at the positions of 1,581 and 1,582 are not 3'-AG but -C. Therefore, in Fig.3 on page 7,324 the open reading frame of NP gene (OP-1) should be assigned to 1,551 bases instead of 1,572 bases, in Fig.4 on page 7,325 the length of the predicted amino acid sequence of OP-1 should be changed to 517 amino acid residues instead of 524, and the sequence at the positions from 488 to 524, -SRE TARDSATHGD EGRNNGVDYD EDDDTAAVAG VGGI (524) should be replaced by -AER RQEILQPMEM KAAITVSIMT KMTIPQQ (517). This resulted in the change of the molecular weight of NP protein from 57,017 to 56,697. The correction gives little influence on the conclusions described in the communications.