

**Supplementary Figure 1. Amino acid sequence alignment of the H proteins of various CDV and MV strains.** The amino acid sequences at positions 503–560 of the CDV H proteins and positions 507–564 of the MV H proteins are aligned. Dots indicate the identical amino acid residues to those of the CDV CYN07-dV H protein. The box indicates the amino acid at position 541 of the CDV H protein and position 545 of the MV H protein. The DDBJ/GenBank accession numbers are provided with the CDV and MV strain names.