

**Supplemental Information**

**for**

**Genetic determinants restricting the reassortment of heterologous**

**NSP2 genes into the simian rotavirus SA11 genome**

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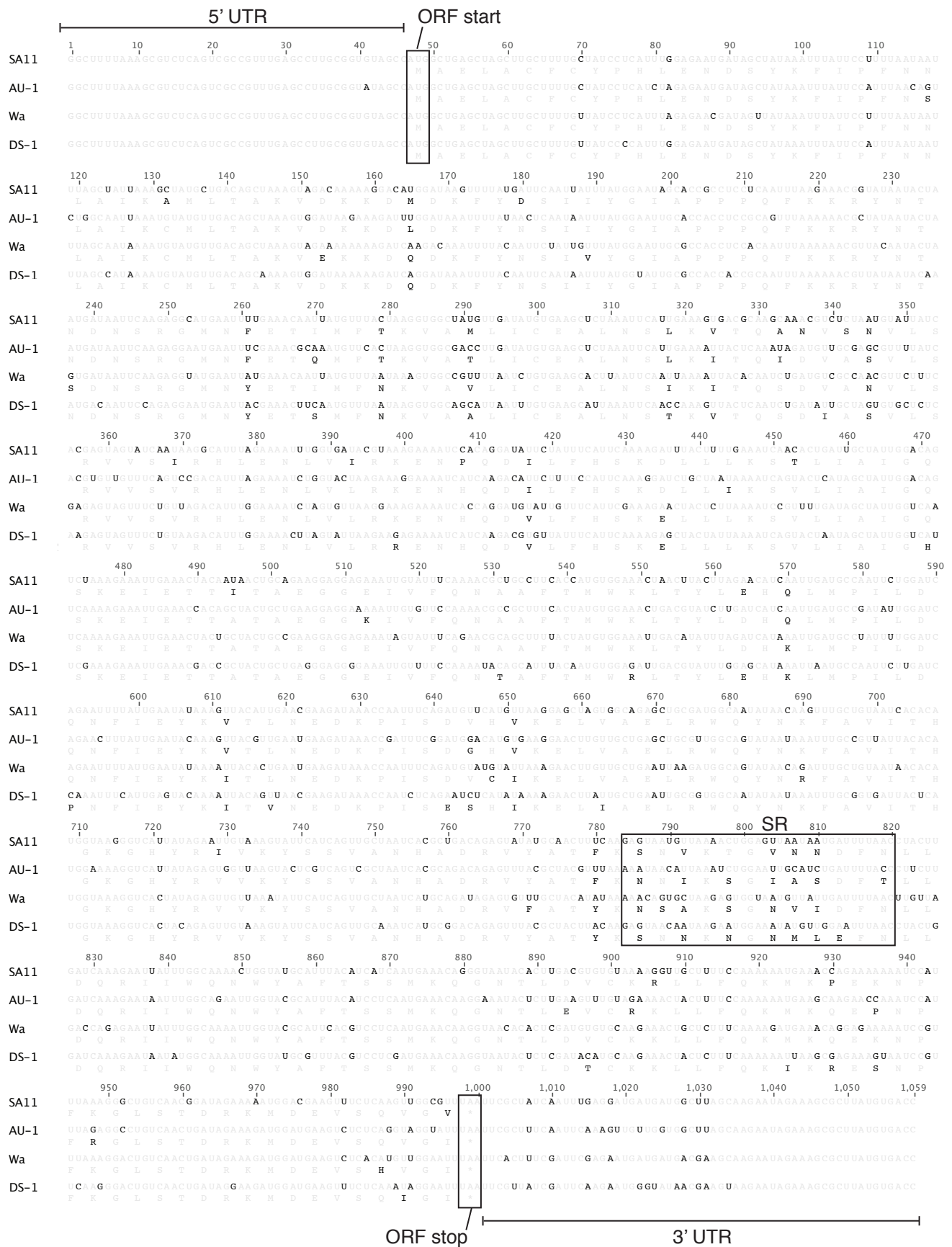
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**Fig. S1. Alignment of NSP2 genes/proteins for RV strains SA11, AU-1, Wa, and DS-1.** Strain names are listed to the left of each nucleotide sequence and position numbers are listed on the top. Deduced amino acid sequences for NSP2 proteins are shown underneath the corresponding nucleotide sequence. Conserved residues are shown in grey and variable residues are shown in black. The locations of the UTRs are shown with brackets and labeled. The ORF start/stop codons and SR are outlined with black boxes and labeled.