

**S2 Table. Primers used in conventional RT-PCR and Sanger sequencing.**

Gene	Primer	Polarity <sup>a</sup>	Sequence (5'-3')	Nucleotide Position <sup>b</sup> (nt)	Reference
<b>VP7</b>	Beg9	+	GGCTTAAAAGAGAGAATTCCGTCTGG	1-28	[1]
	End9	-	GGTCACATCATACAATTCTAATCTAAG	1062-1036	[1]
	Mid1r	-	ATTCCTAACGTTGTGTATT	662-643	[2]
	Mid2	+	GATTGATATTGAATGAATGG	517-537	[2]
	VP7-1`F	+	AAAGGATGCCAACAGGATCAGT	373-395	[3]
<b>VP4</b>	con3	+	TGGCTTCGCTCATTATAGACA	11-32	[4]
	con2	-	ATTCGGACCATTATAACC	887-868	[4]
<b>VP6</b>	VP6-F	+	GGCTTTAACGAAGTCTTC	1-20	[5]
	VP6-R	-	GGTCACATCCTCTCACT	1356-1339	[5]
<b>NSP2</b>	NSP2-F	+	GGCTTTAACCGTCTCAGTC	1-21	[6]
	NSP2-R	-	GGTCACATAAGCGCTTCTATT	1058-1036	[6]

<sup>a</sup> +, Forward; -, Reverse.

<sup>b</sup> The positions correspond to VP7 (K02033), VP4 (FJ423116), VP6 (FJ423150) , NSP2 (FJ423152) of reference Wa strain.

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