

Supplementary Table 3. Estimates of evolutionary divergence rate between the amino acid (A) and nucleotide (B) sequences of the VP1 of the assembled genomes. The rate variation among sites was modelled with a gamma distribution (shape parameter = 1). The actual number of differences are provided in the parentheses.

A	BG 1 VP1	BG 2 VP1	BG 3 VP1	BG 4 VP1	BG 5 VP1	BG 6 VP1
BG 1 VP1	0 (0)	0.00118 (1)	0 (0)	0.00118 (1)	0.00237 (2)	0.00237 (2)
BG 2 VP1		0 (0)	0.00118 (1)	0 (0)	0.00237 (2)	0.00237 (2)
BG 3 VP1			0 (0)	0.00118 (1)	0.00237 (2)	0.00237 (2)
BG 4 VP1				0 (0)	0.00237 (2)	0.00237 (2)
BG 5 VP1					0 (0)	0 (0)
BG 6 VP1						0 (0)

B	BG 1 VP1	BG 2 VP1	BG 3 VP1	BG 4 VP1	BG 5 VP1	BG 6 VP1
BG 1 VP1	0 (0)	0.00158 (4)	0.000395 (1)	0.00158 (4)	0.00592 (15)	0.00553 (14)
BG 2 VP1		0 (0)	0.00118 (3)	0 (0)	0.00474 (12)	0.00434 (11)
BG 3 VP1			0 (0)	0.00118 (3)	0.00553 (14)	0.00513 (13)
BG 4 VP1				0 (0)	0.00474 (12)	0.00434 (11)
BG 5 VP1					0 (0)	0.000395 (1)
BG 6 VP1						0 (0)