

Electronic Supplementary Material

Pathogenic Characterization and Full Length Genome Sequence of a Reassortant Infectious Bursal Disease Virus Newly Isolated in Pakistan

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Table S1 Orientation and position of the virus-specific sequences of the primers.

Name	Sequence (5'-3')	Orientation	Position (nt) ^a
A628U	CCTCAGCTTACCCACATC	Sense	A: 628-645
A1540L	CCTTCCCAATTGCATGG	Antisense	A: 1540-1557
AU	GGATACGATCGGTCTGACCCCGGGGGAGTC	Sense	A: 1-30
A1542L	GTAGTCTACACCTTCCCAATTGCAT	Antisense	A: 1542-1567
A1421U2	GACCTCAACTCTCCCTGAAGATTGCAGGAG	Sense	A: 1421-1451
AL2	GGGGACCCGCGAACGGATCCAATTTGGGATGTTG	Antisense	A: 3227-3260
BU	GGATACGATGGGTCTGACCCTCTGGGA	Sense	B: 1-27
B1344L	TCTAGGTCAATTGAGTACCAC	Antisense	B: 1344-1364
B1344U	GTGGTACTCAATTGACCTAGA	Sense	B: 1344-1364
BL	ttGGGGGCCCCCGCAGGCGAAGGCCGGGGATA	Antisense	B: 2798-2827

^a The positions where the primers bind are in accordance with the published sequence of IBDV strain Gx (GenBank accession nos: AY444873 and AY705393).