

Genome segment ISAV strain

Sequence is in 5' to 3' orientation

1 10 20 30 40

Segment 1	NBISA01	(HQ011270)	AGCTAAGA
Segment 1	ADL-ISAV-07	(HQ011262)	AGCTAAGA
Segment 2	NBISA01	(HQ011271)	AGCAAAGAACGCTCTTTAATAACC
Segment 2	ADL-ISAV-07	(HQ011263)	AGCAAAGAACGCTCTTTAATAACC
Segment 3	NBISA01	(HQ011272)	AGCAAAGATTGCTCAAATCCCAAAAATAATACAGAAAACGTATAAGAG
Segment 3	ADL-ISAV-07	(HQ011264)	AGCAAAGATTGCTCAAATCCCAAAAATAATACAGAAAACGTATAAGAG
Segment 4	NBISA01	(HQ011273)	AGCTAAGATTGGCTATTTTCAAG
Segment 4	ADL-ISAV-07	(HQ011265)	AGCTAAGATTGGCT-GTTTCAAG
Segment 5	NBISA01	(HQ011274)	AGTTAAAG
Segment 5	ADL-ISAV-07	(HQ011266)	AGTTAAAG
Segment 6	NBISA01	(HQ011275)	AGCAAAG
Segment 6	ADL-ISAV-07	(HQ011267)	--CAAAG
Segment 7	NBISA01	(HQ011276)	AGCTAAGATTCTCCTTCTACA
Segment 7	ADL-ISAV-07	(HQ011268)	AGCTAAGATTCTCCTTCTACA
Segment 8	NBISA01	(HQ011277)	AGCAAAGATTGGCTATTTACC
Segment 8	ADL-ISAV-07	(HQ011269)	AGCAAAGGTTGGCTATCTACC