

Segment	Type (SUBTYPE)	NB of isolates	Position (H3 and N2 numbering)	Wild	Mutant	Antiviral resistance			High patho- genicity	Hum. adapt.	Ref.	
						Neuraminidase inhibitors						Adam.
						Ose.	Zan.	Per.				
NA	A(N2), B	12	119	E	V	X					38, 40, 42, 43	
NA	A(N2), B	1	119	E	A	X	X				42, 43	
NA	A(N2), B	3	119	E	G		X				42, 43	
NA	A(N2), B	7	119	E	D		X	X			42, 43	
NA	A(N1)	415	274	H	Y	X		X			34-38, 41, 43, 44	
NA	A(N1,N2)	0	294	N	S	X					33	
NA	A(N2), B	2	292	R	K	X	X	X			39, 42, 43	
NA	B	0	152	R	K	X	X	X			36-38, 42, 43	
NA	B	3	198	D	N/E	X	X				38, 43, 44	
NA	B	1	222	I	T	X	X				44	
NA	A(N2)	6	222	I	V	X	X				44	
M2	A	20	26	L	F				X		46	
M2	A	130	27	V	A				X		45, 46	
M2	A	8	30	A	T				X		18, 46	
M2	A	1433	31	S	N				X		18, 45, 46	
M2	A	0	34	G	E				X		46	
HA	A	2030	HA cleavage site		≥ 5 R or K					+++	24, 25	
NP	A	92	319	N	K					++	31	
PB2	A	3181	627	E	K					++	29	
PB2	A	177	701	D	N					++	29, 30, 31	
NS1	A	97	92	D	E					+ (M)	27, 26	
NS1	A	1240	80-84	XXXXX	-					+ (M)	27	
PB1-F2	A	1416	66	N	S					++	28	

Table 1: List of mutations used to compute antiviral resistance and putative human adaptation. Ose.: Oseltamivir. Zan.: Zanamivir. Per.: Peramivir. Adam.: Adamantanes. Hum. adapt.: Human adaptation. Ref.; bibliographic reference number. High pathogenicity: could contribute, alone or in association with other mutations, to a higher pathogenicity in avians and/or mammals. +++: high level of confidence; ++: medium level of confidence; +: low level of confidence, or in correlation with other mutations; (M): mammals only.

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