

TABLE 1. Distribution of Influenza Subclade Diversity, by Influenza A Subtype and B/Victoria Lineage Virus, U.S. Department of War, 2025–2026

Subtype, Lineage	Clade	Subclade	No.	% of Total Subtype	Defining Substitutions ^a
A(H1N1)pdm09	5a.2a	C.1.9	3	2.2	K169Q
		C.1.9.3	7	5.1	S83P, K169Q
	5a.2a.1	D.3.1 ^b	72	52.2	I372V, I460T, V520A
		D.3.1.1	56	40.6	R113K, A139D, E283K, K302E
A(H3N2)	2a.3a.1	J.2 ^c	14	1.3	K276E
		J.2.2	6	4.3	T65K, S124N
		J.2.3	7	0.7	S378N
		J.2.4	39	3.7	T135K
		K ^d	975	93.7	K2N, Q173R
B/Victoria	V1A.3a.2	C.3	16	13.2	A154E, S208P, S255P
		C.3.1 ^e	87	71.9	D197N
		C.5.1	2	1.7	E128K, E183K
		C.5.6	11	9.1	D128E
		C.5.6.1	4	3.3	T37I, P241Q
		C.5.7	1	0.8	E128G, E183K

Abbreviation: No., number.

^a All substitutions reference changes in hemagglutinin subunit 1 (HA1).

^b Subclade of selected 2026-2027 Northern Hemisphere A(H1N1)pdm09 strain A/Missouri/11/2025 (egg, cell).

^c Subclade of previous 2025-2026 Northern Hemisphere A(H3N2) strains A/Croatia/10136RV/2023 (egg) and A/District of Columbia/27/2023 (cell).

^d Subclade of selected 2026-2027 Northern Hemisphere A(H3N2) strains A/Darwin/1454/2025 (egg) and A/Darwin/1415/2025 (cell).

^e Subclade of selected 2026-2027 Northern Hemisphere B/Victoria strains B/Tokyo/EIS13-175/2025 (egg) and B/Pennsylvania/14/2025 (cell).