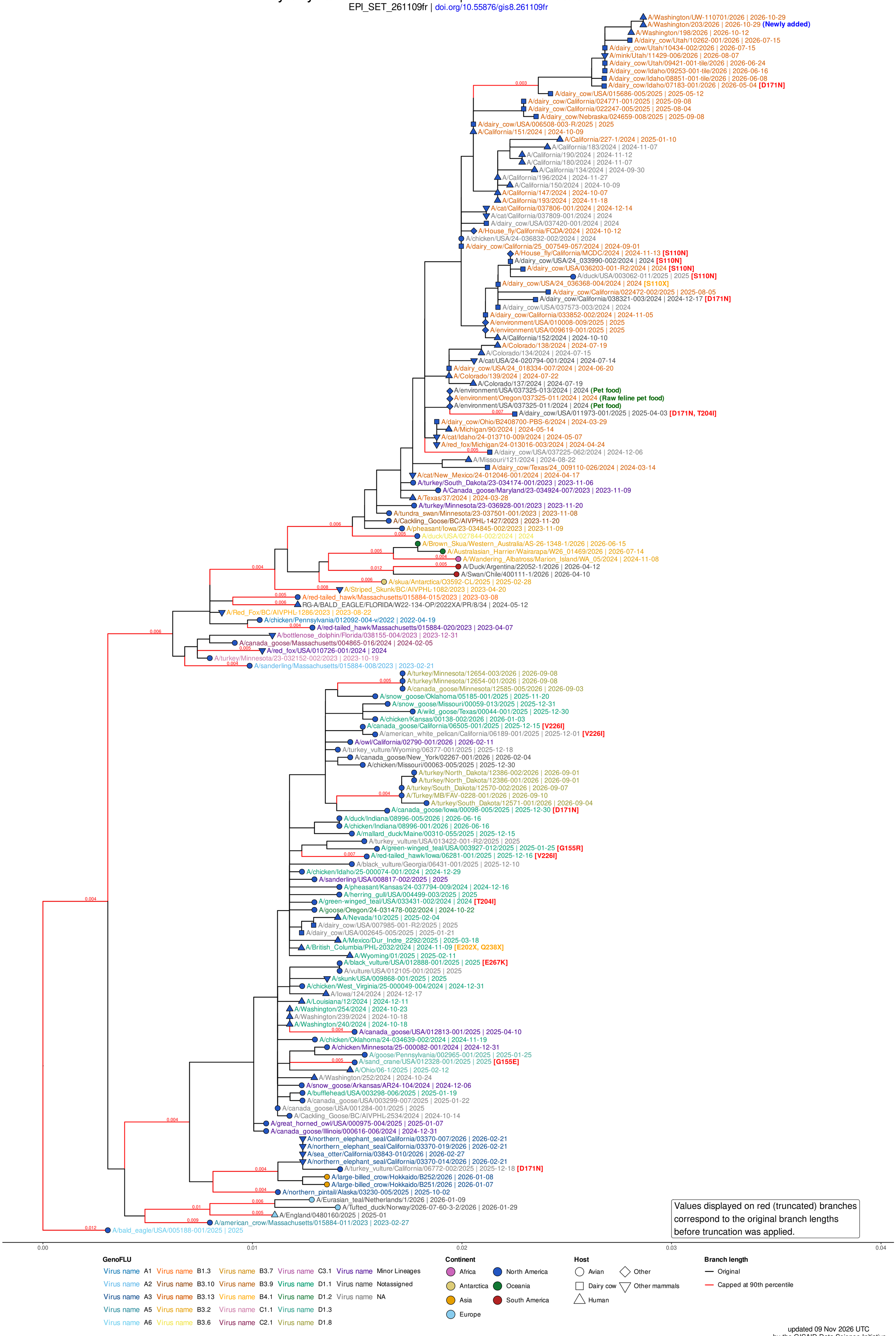
Phylodynamics of H5N1 HA | North America outbreak

EPI_SET_261109fr | doi.org/10.55876/gis8.261109fr



Values displayed on red (truncated) branches correspond to the original branch lengths before truncation was applied.

GenoFLU

Virus name	A1	Virus name	B1.3	Virus name	B3.7	Virus name	C3.1	Virus name	Minor Lineages
Virus name	A2	Virus name	B3.10	Virus name	B3.9	Virus name	D1.1	Virus name	Notassigned
Virus name	A3	Virus name	B3.13	Virus name	B4.1	Virus name	D1.2	Virus name	NA
Virus name	A5	Virus name	B3.2	Virus name	C1.1	Virus name	D1.3		
Virus name	A6	Virus name	B3.6	Virus name	C2.1	Virus name	D1.8		

Continent

Africa · North America · Antarctica · Oceania · Asia · South America · Europe

Host

Avian · Other · Dairy cow · Other mammals · Human

Branch length

Original · Capped at 90th percentile

updated 09 Nov 2026 UTC
by the GISAID Data Science Initiative