Probable extinction of influenza B/Yamagata and its public health implications: a systematic literature review and assessment of global surveillance databases

CONTEXT

Influenza viruses of the B/Yamagata lineage have only been detected in a few countries globally since the start of the COVID-19 pandemic in March 2020

In this study, the circulation of B/Yamagata influenza viruses from 2020 onwards was investigated, to understand whether these viruses are extinct and consider the public health implications of the findings¹

STUDY AIM

To conduct an updated systematic review of the evidence pertaining to global B/Yamagata influenza virus circulation, building on previous assessments by including multiple data sources¹

METHODS

The study methodology, conducted in January 2024, utilised 3 different types of data sources to search for B/Yamagata influenza viruses:1



A systematic review of the literature published between 1 January 2020 – 2 January 2024 using the PubMed and Embase databases



A search of **four influenza surveillance databases** –
FluNet, GISHN, GISAID and
GenBank, for data reported
during the period 2020–2023



A review of the webpages of national respiratory virus surveillance systems of 21 countries* for data reported during the period 2020–2023

*Argentina, Australia, Brazil, Canada, China, England, France, Germany, Hong Kong, Israel, Italy, Japan, Mexico, New Zealand, Portugal, Russia, South Africa, Spain, the Netherlands, Türkiye, US.

KEY FINDINGS¹

Literature review

The literature search returned **5152** non-duplicate articles of which **43** were suitable for inclusion

In studies with observation periods from May 2020 through July 2023, **B/Yamagata detections were reported in**:

only 4 countries globally

<2 in 1000 of all influenza B cases reported

Influenza database findings

FluNet

From **2,376,287** reported influenza detections between 2020 and 2023:

435,250 (18.3%) were influenza type B

Among these **influenza type B** detections:

297,915 (68.4%) were not characterised **137,015 (31.5%)** were B/Victoria lineage

320 (<0.1%) were of B/Yamagata lineage

The average number of influenza B/ Yamagata detections per 100,000 processed specimens dropped from:

14.9 in weeks 1–17 of **2020**

to **<0.10** in **2022** and **2023**

GISHN

From **9,204** influenza detections reported to GIHSN between 2020 and 2023:

2168 (23.6%) were influenza type B

Among **1030 influenza type B** viruses that were characterised:

Seven (0.7%) were B/Yamagata, all detected during 2020

No B/Yamagata influenza detections were reported from 2021 onwards

GISAID and GenBank

No sequences of wild-type viruses of the B/Yamagata lineage with a collection date after March 2020 were found in the GISAID and GenBank databases

National respiratory virus surveillance system webpages

Twelve B/Yamagata detections were reported after week 35 of 2020:

1 in Australia (week 37–38 of 2020)

10 in the US (weeks 44–49 of 2021)

1 in Spain (in 2023; with a sequence identical to the live attenuated vaccine strain)

The current protracted and nearly undetectable viral circulation of the B/Yamagata lineage provides enough evidence to suggest that these viruses are on the verge of extinction, if they are not extinct already¹

On March 5, 2024, the FDA's Vaccines and Related Biological Products Advisory Committee

an influenza A(H1N1), an A(H3N2) and a B/Victoria-lineage vaccine virus²

To download a copy of this infographic visit the Nivel FluCov website:

FluCov: Influenza-COVID-19, understanding and communicating the impact of COVID-19 on influenza activity | Nivel

(VRBPAC) recommended that all 2024–2025 US flu vaccines be trivalent vaccines and include

2025 | CDC. Accessed July 2024.