

Questions and Answers

Recommended composition of influenza virus vaccines for use in the northern hemisphere 2026-2027 influenza season and development of candidate vaccine viruses for pandemic preparedness

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1. What is the purpose of WHO recommendations on the composition of influenza virus vaccines?.....	2
2. What is the Global Influenza Surveillance and Response System (GISRS)?.....	2
3. How are influenza virus vaccine recommendations made?.....	3
• Surveillance data:	3
• Antigenic characterization of viruses:	3
• Human serology studies with influenza virus vaccines:.....	3
• Genetic characterization of viruses:.....	3
• Virus fitness forecasting:.....	3
• Antiviral susceptibility:	3
• Vaccine effectiveness:	3
• Availability of CVVs:	3
4. What are candidate vaccine viruses (CVVs)?.....	4
5. Which viruses are recommended by WHO to be included in influenza virus vaccines for use in the 2026-27 northern hemisphere influenza season?.....	4
6. Are the vaccine viruses in this recommendation different from those in the previous northern and southern hemisphere recommendations?	4
7. What is the seasonal influenza A(H3N2) “subclade K” virus?	5
8. What does the term “-like virus” mean in the vaccine composition recommendation?.....	5
9. Why are different viruses sometimes recommended for egg- and cell-based vaccines?..	5
10. Who recommends, approves, advises on influenza virus vaccine composition and use?.....	5
11. What vaccine formulation (i.e., recommendation for northern or southern hemisphere influenza season) should countries in tropical and subtropical regions consider for use in vaccination programmes?.....	6
12. Which CVVs are available for use in influenza virus vaccines?	6
13. Why and how does GISRS continue to update CVVs for pandemic preparedness purposes?.....	6
14. What is WHO’s guidance on the use of influenza A(H5) virus vaccines?	7

1. What is the purpose of WHO recommendations on the composition of influenza virus vaccines?

These WHO recommendations provide a guide to national public health and regulatory authorities, and vaccine manufacturers for the development and production of influenza virus vaccines for the next influenza season and for pandemic preparedness. In contrast to many other vaccines, the viruses in seasonal influenza vaccines need to be evaluated and updated regularly because circulating influenza viruses evolve continuously and vaccines may need to be updated to remain effective. Recommendations are usually made in February for the following influenza season in the northern hemisphere and in September for the following influenza season in the southern hemisphere. The recommendation dates are chosen to provide approximately 6-8 months for the production, regulatory approval and distribution of the manufactured vaccines.

For pandemic preparedness, zoonotic influenza candidate vaccine virus (CVV) selection and development are considered at least twice a year. The decisions are based on continuous surveillance for zoonotic influenza events (human infection with an influenza virus normally restricted to a non-human host) and influenza virus activity in animals, and virus characterization by Global Influenza Surveillance and Response System (GISRS) and its collaborators e.g., the World Organisation for Animal Health, founded as OIE (WOAH), Food and Agriculture Organization of the United Nations (FAO), and the WOAH/FAO Network of Expertise on Animal Influenza (OFFLU).

2. What is the Global Influenza Surveillance and Response System (GISRS)?

GISRS is a global system of public health institutions coordinated by WHO, currently consisting of 165 institutions, including National Influenza Centres (NICs), WHO Collaborating Centres for Influenza (CCs), WHO Essential Regulatory Laboratories (ERLs) and WHO H5 Reference Laboratories, in 135 WHO Member States. The GISRS functions year-round under WHO [Terms of Reference](#), together with partner laboratories and its collaborators, sharing surveillance findings and virus materials from human and animal health sectors in a timely fashion to inform risk assessment and mitigation measures, and serves as the global alert mechanism for emerging respiratory threats.

GISRS monitors the evolution of influenza viruses of public health importance, including seasonal, zoonotic and potential pandemic viruses, and recommends and implements risk assessment and response measures. Virus characterizations, combined with other available epidemiological and disease information, form the evidence base for public health decisions on epidemic response and pandemic preparedness including selection and development of candidate vaccine viruses (CVVs) for seasonal and zoonotic influenza. GISRS also provides guidance to countries and support for activities such as training, risk assessment, outbreak response, development of diagnostic tests, testing for antiviral drug resistance and scientific interpretation of important findings.

GISRS has been expanding its scope to include other respiratory viruses. Respiratory syncytial virus (RSV) and severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) have been included in the GISRS surveillance platform since 2015 and 2020, respectively. The broader application and value of GISRS have been increasingly acknowledged by the world.

Further information about GISRS is available on the [WHO website](#).

3. How are influenza virus vaccine recommendations made?

Data and information from the GISRS network, which includes NICs, WHO CCs, WHO ERLs and WHO H5 Reference Laboratories, its collaborators, and other sources are used to make vaccine virus recommendations. This includes:

- ***Surveillance data:***

Virus surveillance data from the GISRS network, complemented with epidemiologic and clinical findings, inform the vaccine virus selection process.

- ***Antigenic characterization of viruses:***

GISRS laboratories, in particular WHO CCs, use post-infection ferret antisera and pooled post-vaccination human sera to evaluate antibody reactivity with the surface proteins (antigens) of circulating and vaccine influenza viruses. Antigenic cartography is used to visualize relatedness of viruses based on the data provided by WHO CCs.

- ***Human serology studies with influenza virus vaccines:***

WHO CCs and WHO ERLs test how well antibodies from vaccinated people react with recently circulating influenza viruses.

- ***Genetic characterization of viruses:***

GISRS laboratories conduct gene sequencing to compare the sequences of circulating influenza viruses with those of vaccine viruses to identify genetic changes that might influence protection conferred by a given influenza vaccine.

- ***Virus fitness forecasting:***

Virus fitness relates to the likelihood of any emerging groups of viruses becoming more prevalent. Information from modelling studies, based on genetic sequences available in databases and antigenic information provided by the WHO CCs is considered.

- ***Antiviral susceptibility:***

GISRS laboratories analyse influenza viruses to determine if they remain susceptible to the antiviral drugs approved for treatment of influenza infections. This information is taken into consideration when specific viruses are selected as CVVs.

- ***Vaccine effectiveness:***

The Global Influenza Vaccine Effectiveness (GIVE) Collaboration, made up of many different studies conducted in countries in both the northern and southern hemispheres, provides information on vaccine performance in previous influenza seasons and interim reports on the current season.

- ***Availability of CVVs:***

CVVs are essential for production of vaccines in a timely manner for the next influenza season. Most vaccines produced globally use egg-based manufacturing processes which require CVVs that replicate well in eggs while cell-based vaccines require CVVs that replicate well in cell culture. Separate recommendations are made for egg- and cell-base

CVVs because of the differing replication and manufacturing processes. Influenza virus vaccines comprised of recombinant protein influenza virus antigens do not require CVVs for manufacturing.

These data, and other findings made available by GISRS, are evaluated during WHO consultations held in February and September of each year. The consultations include experts from WHO CCs, WHO ERLs, WHO H5 Reference Laboratories, NICs, OFFLU, academic institutions, and other national and regional institutions.

4. What are candidate vaccine viruses (CVVs)?

Haemagglutinin (HA) is the primary antigen in seasonal influenza vaccines. A CVV is a virus prepared for potential use in vaccine manufacturing that possesses an HA which WHO CCs have determined to be antigenically similar to the virus that has been recommended for use in vaccines.

5. Which viruses are recommended by WHO to be included in influenza virus vaccines for use in the 2026-27 northern hemisphere influenza season?

WHO recommends that vaccines for use in the 2026-27 northern hemisphere influenza season contain the following:

Egg-based vaccines

- an A/Missouri/11/2025 (H1N1)pdm09-like virus;
- an A/Darwin/1454/2025 (H3N2)-like virus; and
- a B/Tokyo/EIS13-175/2025 (B/Victoria lineage)-like virus.

Cell culture-, recombinant protein- or nucleic acid-based vaccines

- an A/Missouri/11/2025 (H1N1)pdm09-like virus;
- an A/Darwin/1415/2025(H3N2)-like virus; and
- a B/Pennsylvania/14/2025 (B/Victoria lineage)-like virus.

6. Are the vaccine viruses in this recommendation different from those in the previous northern and southern hemisphere recommendations?

The recommended composition of influenza vaccines for the 2026-27 northern hemisphere is different from those viruses recommended for the 2025-26 northern hemisphere and the 2026 southern hemisphere vaccines.

The A(H1N1)pdm09, A(H3N2), and B/Victoria components differ from the previous 2025-26 northern hemisphere vaccine composition, while the A(H3N2) and B/Victoria components differ from the 2026 southern hemisphere vaccine composition.

Previous and present WHO influenza virus vaccine composition recommendations can be found on the WHO Global Influenza Programme [website](#).

7. What is the seasonal influenza A(H3N2) “subclade K” virus?

Subclade K viruses are human seasonal A(H3N2) influenza viruses that emerged from the J.2.4 subclade (included in the southern hemisphere 2026 influenza vaccine composition recommendation). Being seasonal influenza viruses, they are not subject to notification requirements under the [International Health Regulations \(IHR\)](#). In the September 2025 [WHO vaccine composition recommendation for the 2026 southern hemisphere influenza season](#), a group of subclade J.2.4. viruses was identified with notable additional HA substitutions S144N (a potential addition of an N-glycosylation site), N158D, I160K and Q173R. This group of viruses was classified as subclade J.2.4.1 in October and then aliased as subclade K using the nomenclature defined by HA substitutions K2N, S144N, N158D, I160K, Q173R, T328A and S378N¹.

8. What does the term “-like virus” mean in the vaccine composition recommendation?

The influenza vaccine viruses recommended for a given season are selected because they represent the antigenic groups of viruses expected to circulate widely during the forthcoming influenza season. Often, more than one candidate vaccine virus (CVV) is available. These CVVs contain haemagglutinin (HA) antigens derived from other viruses that are antigenically similar to the recommended vaccine virus. The term “-like virus” is therefore used to indicate that these antigenically similar CVVs may also be used for vaccine manufacturing.

9. Why are different viruses sometimes recommended for egg- and cell-based vaccines?

Influenza viruses may not replicate equally well in the egg- and cell-based vaccine production systems. Influenza viruses may undergo changes when they are grown in eggs, and some egg-adapted changes may result in antigenic changes that reduce recognition of sera raised to egg grown viruses to cell-grown viruses. The WHO influenza vaccine composition recommendations select the best candidates for each production system and take into consideration the egg-adapted changes in available egg isolates. Therefore, different viruses with similar antigenic properties along with their genetic sequences are sometimes recommended for different production systems.

10. Who recommends, approves, and advises on influenza virus vaccine composition and use?

The WHO influenza vaccine composition advisory committee recommends which viruses should be used to make influenza virus vaccines. WHO publishes and updates a [list](#) of

¹ Neher RA, Huddleston J, Bedford T, Lewis NS, Harvey R, Galiano M, et al. Nomenclature for Tracking of Genetic Variation of Seasonal Influenza Viruses. *Influenza Other Respir Viruses*. 2026;20(2):e70230. <https://doi.org/10.1111/irv.70230>. <https://www.ncbi.nlm.nih.gov/pubmed/41688063>.

recommended CVVs and sequence accession numbers. It is the responsibility of the vaccine manufacturers to source the appropriate CVVs or protein sequences and to obtain approval for their use from the relevant regulatory agencies.

Following the global vaccine composition recommendation by WHO, national or regional authorities approve the composition and formulation of vaccines and develop policy for the use of influenza virus vaccines in their jurisdictions.

11. What vaccine formulation (i.e., recommendation for northern or southern hemisphere influenza season) should countries in tropical and subtropical regions consider for use in vaccination programmes?

WHO has developed [guidance](#) to support countries in tropical and subtropical regions in choosing between the northern and southern hemisphere formulations. These countries should consider their epidemiologic and virologic surveillance data in selecting which vaccine formulation to use and deciding when to start vaccination.

12. Which CVVs are available for use in influenza virus vaccines?

The WHO recommended CVVs for vaccine development and production for the 2026-27 northern hemisphere influenza season are listed on the [WHO website](#).

The available CVVs, including those for pandemic preparedness purposes ([zoonotic viruses](#)), and corresponding potency test reagents are posted by type/subtype and updated on the [WHO website](#).

13. Why and how does GISRS continue to update CVVs for pandemic preparedness purposes?

Animal influenza viruses circulate widely in some animals and transmit sporadically to humans, resulting in zoonotic infections. Human infections with influenza viruses from birds, pigs and other mammals including dairy cattle continue to be detected, and these viruses remain a public health threat. As such, GISRS and its collaborators continue to conduct surveillance and risk assessments² of zoonotic influenza and update CVVs for pandemic preparedness purposes.

GISRS, in collaboration with animal health partners, continuously analyses a range of zoonotic and potentially pandemic influenza viruses as they emerge and evolve. Sharing of virus materials with a WHO CC is crucial for further characterization and CVV development and analysis strengthens pandemic preparedness and highlights the importance of the One Health approach through cross sectoral cooperation. Data are reviewed at least twice a year. Influenza viruses with potential public health significance and not well covered by existing CVVs are considered as a priority for new CVV development.

² Monthly and ad-hoc risk assessment of zoonotic influenza can be found on the WHO Global Influenza Programme website at: <https://www.who.int/teams/global-influenza-programme/avian-influenza/monthly-risk-assessment-summary>.

CVVs are developed by classical reassortment or by recombinant DNA technology, and undergo extensive quality and safety testing to ensure the CVVs are phenotypically attenuated and antigenically similar to the selected wild-type virus. An inventory of CVVs and associated potency reagents is maintained and updated for various pandemic preparedness purposes, e.g., vaccine development to conduct clinical trials, and other pandemic preparedness activities. The use of these CVVs for vaccine development and production should be based on the assessment of the public health risk and the needs in consultation with national and/or regional regulatory and public health authorities.

A list of available pre-pandemic CVVs can be found on the [WHO website](#).

14. What is WHO's guidance on the use of influenza A(H5) virus vaccines?

In 2008, WHO published its first guidance on the subject, "[Options for the use of human H5N1 influenza vaccines and the WHO H5N1 vaccine stockpile](#)", which was updated and replaced in 2025 by "[Considerations for use of avian influenza A\(H5\) vaccines during the interpandemic and emergence periods](#)."

The WHO Scientific Advisory Group of Experts on Immunization (SAGE) made its recommendations on the "use of licensed human H5N1 influenza vaccines in the interpandemic period" in [2009](#), and reconfirmed its recommendations in [2013](#). In 2025, SAGE issued [updated recommendations](#) on the "use of licensed human A(H5) vaccines for the interpandemic and emergence periods."

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