

From Data to Action: Understanding how WHO's Global Surveillance on Coronaviruses informs Response

Introduction

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Team Lead, WHO Global COVID-19 and other Coronaviruses Programme

5 November 2025





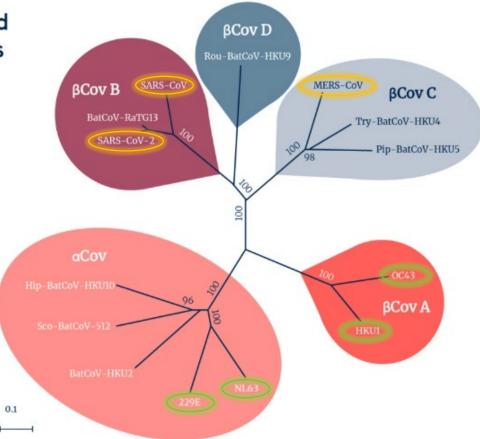
Coronaviruses

The Coronavirus family includes hundreds of viruses, divided into four groups:

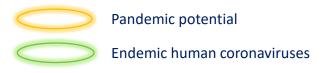
- Alphacoronaviruses
- Betacoronaviruses
- Gammacoronaviruses
- Deltacoronaviruses

Most human diseases are caused by Alpha- and Betacoronaviruses

Phylogenetic Tree and Genetic Relationships of Different Coronaviruses





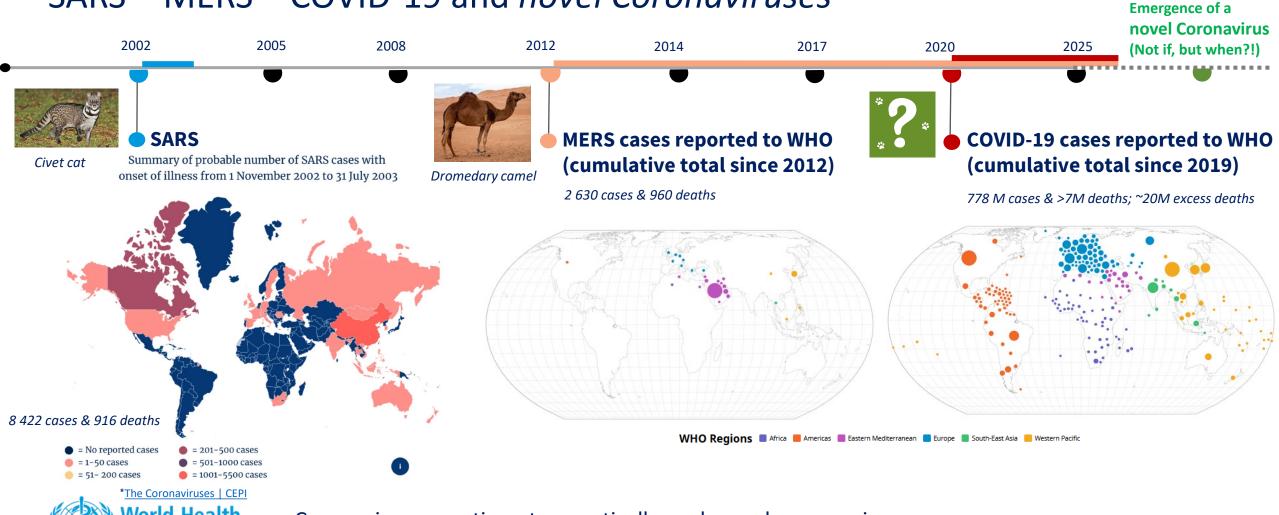








Past, present and future zoonotic coronavirus threats: SARS – MERS – COVID-19 and *novel Coronaviruses*



Coronaviruses continue to genetically evolve and can acquire new features, such as human infectiousness, increased virulence, etc.



The strategic direction of coronavirus disease threat management efforts continues to evolve





From Emergency Response to Long-Term COVID-19

<u>Disease Management:</u> Sustaining Gains Made

During the COVID-19 Pandemic

Goal: End the emergency phase of the COVID-19 pandemic in all countries and shift from emergency response to sustainable comprehensive management of COVID-19 within broader disease prevention and control programmes.





A GLANCE



Strategic and operational plan for coronavirus disease threat management

Advancing sustained, integrated coronavirus disease threat management: 2025-2030

2025 - onwards

Shift into the programmatic management of coronavirus disease threats, incl. COVID-19, MERS, etc.

EMERGENCIES

WHO's Global Coronavirus Programme is organized around the 5 Cs of the HEPR framework and its Strategic Plan has four objectives

<u>5 C's of the Health Emergency Prevention, Preparedness, Response,</u> and Resilience (HEPR) framework



Strategic and operational plan for coronavirus disease threat management Advancing sustained, integrated coronavirus disease threat management: 2025-2030

Strategic objectives

The plan aims to support and guide Member States and the broader global health community to:

- **Sustain** essential, evidence-based COVID-19 and other coronavirus disease threat management activities across core public health capabilities to reduce morbidity, mortality, and socioeconomic disruption, right-sized to burden.
- Integrate coronavirus disease threat management into broader disease prevention and control programmes and systems, across all levels (local, national, regional, global), in particular with other respiratory diseases, like influenza and respiratory syncytial virus (RSV).
- **Enhance** core capabilities as outlined in the HEPR Framework to identify, prioritize, and address operational gaps in coronavirus disease threat management.
- **Generate**, share, and apply evidence to close knowledge gaps and translate research and lessons learned into improved programmes, policies, and evidence-based guidance and control tools.





Surveillance / monitoring⇒ risk evaluation ⇒ decision making

WHO monitors and assesses coronaviruses and evaluates their impact on countermeasures, including vaccines, therapeutics, diagnostics or effectiveness of public health and social measures.



TAG-VE

- Determines which variants warrant further investigations
- Assesses whether variants alter
 - transmission or disease characteristics or
 - performance of vaccines, therapeutics, diagnostics or
 - effectiveness of public health and social measures

Ayşe Açma – Epidemiologist

James Otieno – Bioinformatician/

World Health Genomics

Vaccine

Research, evidence & assessment

R&D Blueprint for epidemics

 COVID-19 Vaccines Research Expert Group

TAG-CO-VAC

- Assesses the impact of SARS-CoV-2 variants on current COVID-19 vaccines
- Determines whether changes to vaccine composition are needed

Bart Haagmans – TAG-VE Chair

Vaccine implementation

Policy

SAGE

 Recommends policies and strategies on vaccine use and immunization programmes through evidence-based assessment

Annelies Wilder-Smith – SAGE Secretariat

Vaccination Programmes

- Global status
- Resources available to support programme strengthening

Shoshanna Goldin – MCM Technical Officer







How WHO monitors the current COVID-19 situation

Ayşe Açma, Epidemiologist

James Richard Otieno, Bioinformatician/Genomic Epidemiologist

5 November 2025

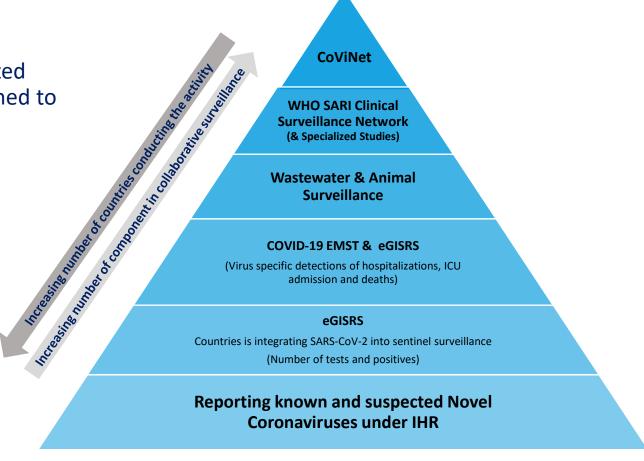




Comprehensive and complementary surveillance approaches for COVID-19 within the Mosaic Framework

Countries can implement multiple surveillance approaches, as coordinated and collaborative systems, well-matched to specific priority objectives











Surveillance types and data flow

- Integrated Sentinel surveillance
- Detailed surveillance
- Genomic surveillance
- | Environmental surveillance
- Event based surveillance
- Vaccine coverage monitoring
- Animal surveillance (Partner Organizations)



Data Triangulated

- Cases & Deaths
- Deaths by Age
- Hospitalizations & ICU
- Test Positivity Rate
- Variants monitored
- Variant distribution
- Vaccination coverage
- Wastewater Sources

Output Types

- Epidemiological Summary
- Maps
- Figures
- Variant Risk Evaluations
- Vaccine Efficacy Plots
- Links to Guidelines & Networks
- Updates and News
- Q&A
- Metadata
- Data Downloadable
- More Resources (SeroTracker)





WHO Global COVID-19 Dashboard

Integration and triangulation of the latest global data and insights on various components of COVID-19 within a unified platform.

COVID-19 Dashboard

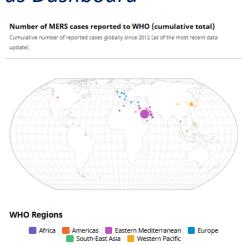
Other Coronavirus Dashboard



Middle East respiratory syndrome coronavirus (MERS-CoV) Dashboard

MERS-CoV Dashboard







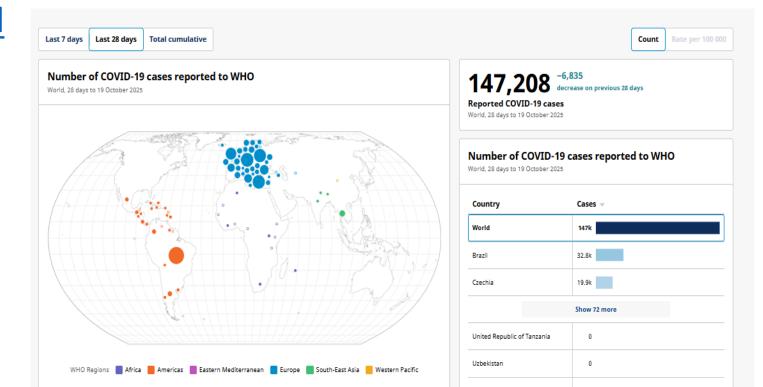
WHO Global ⊕ Language ¾ Hide navigation ‡

re Countries Dashboards



COVID-19 Cases, World

WHO will utilize various sources to continue monitoring the COVID-19 epidemiological situation via the WHO COVID-19 dashboard. If data for certain countries is unavailable in this section, it may indicate that they have either ceased reporting COVID-19 surveillance data to WHO or have integrated the COVID-19 surveillance into existing respiratory disease surveillance. In the latter case, SARS-CoV-2 detections from sentinel and systematic virological surveillance sites in those countries may be found in the Circulation section which also includes information on SARS-CoV-2 variant circulation. This global summary of COVID-19 cases includes data on confirmed cases reported to WHO from the comprehensive COVID-19 case monitoring.



Regional Office Information Sources





erviss.org





respiratory viruses



Covid-19 newsletters

Coronavirus (COVID-19) | WHO | Regional Office for Africa









WHO EMRO - Latest updates



COVID-19 situation reports

Influenza and other respiratory viruses updates

WHO Western Pacific Bi-weekly situation update by date

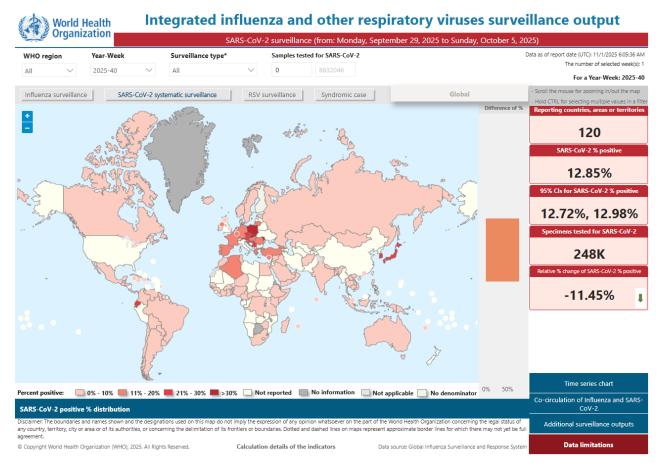
COVID-19 situation reports | WHO Western Pacific





Integrated Surveillance Global Influenza Surveillance and Response System





- Countries are encouraged to integrate SARS-CoV-2 surveillance into existing respiratory disease surveillance systems.
- Currently, 77% of WHO Member States have adopted this more sustainable approach by incorporating SARS-CoV-2 surveillance into influenza sentinel or systematic virological surveillance systems,
- and are reporting data through the expanded GISRS platform.

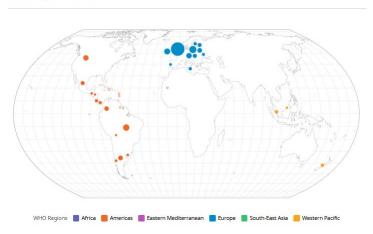




Health impact data (detailed surveillance)

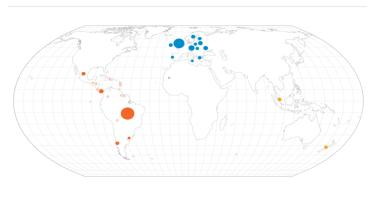
Number of hospitalizations due to SARS-CoV-2 in last 28 days

World, 28 days to 19 October 2025



Number of ICU admissions due to SARS-CoV-2 in last 28 days

World, 28 days to 19 October 2025

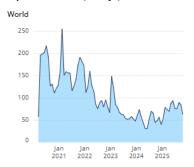


WHO Regions Africa Americas Eastern Mediterranean Europe South-East Asia Western Pacific

World Health Organization

Severity Rate per 1000 Hospitalizations, World

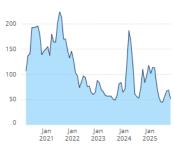
Rate of deaths per 1000 hospitalizations reported to WHO (28 days)



Source: World Health Organization

Rate of ICU admissions per 1000 hospitalizations reported to WHO (28 days)

World

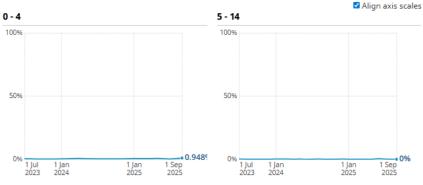


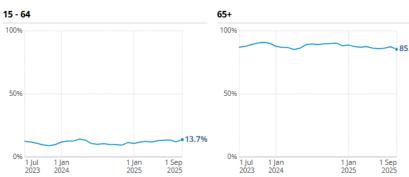
Source: World Health Organization

Data may include corrections and be incomplete for the latest week

Trends in COVID-19 deaths by age, World

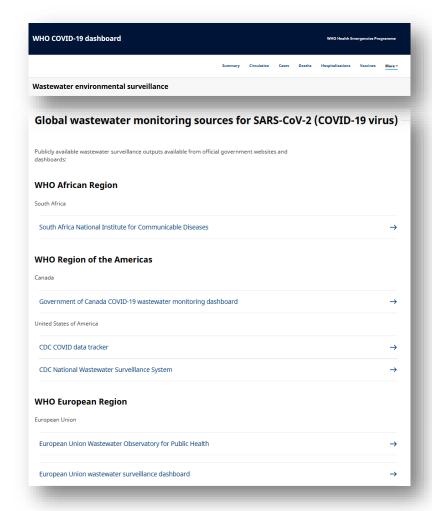
Proportions of deaths reported due to COVID-19 (monthly)

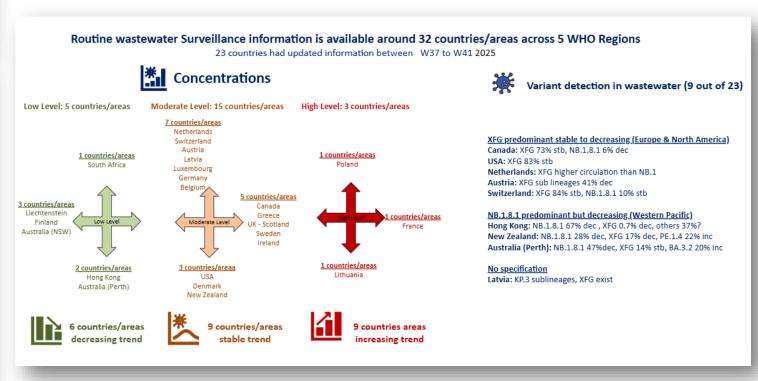






Wastewater Surveillance









Event Based Surveillance

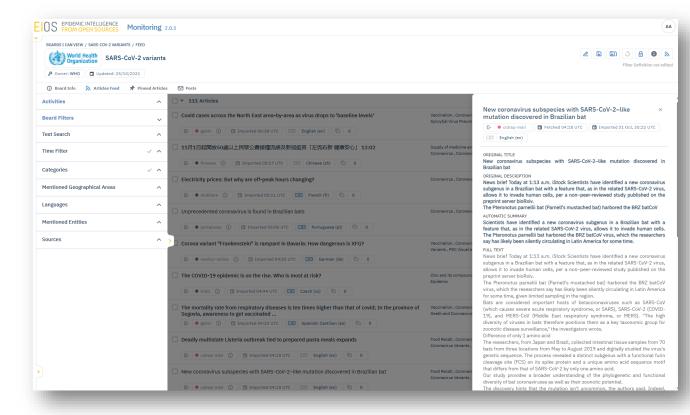
- Official country public health institution websites:
 - dashboards,
 - situation reports,
 - special reports,
 - statements
- 2. Media monitoring (mainly via EIOS)

EIOS is a global WHO-led collaboration that unites public health actors worldwide.

- Articles from selected websites, RSS feeds, social media and other open-source information
- Text mining, algorithms and analytical modules sort and categorize articles
- Adaptable criteria to meet specific surveillance needs

Allows professionals to quickly access multitude of sources based on time period, topic of interest, country/area, language, source type.

Epidemic Intelligence from Open Sources







Animal surveillance via partner organizations

Joint statement on the prioritization of monitoring SARS-CoV-2 infection in wildlife and preventing the formation of animal reservoirs

By Food and Agriculture Organization (FAO), World Organisation for Animal Health (OIE) and World Health Organization (WHO)

7 March 2022 | Statement | Reading time: 3 min (834 words)

<u>Joint statement on the prioritization of monitoring SARS-CoV-2 infection in wildlife</u> and preventing the formation of animal reservoirs

Surveillance and Information Sharing Operational Tool

An operational tool of the Tripartite Zoonoses Guide

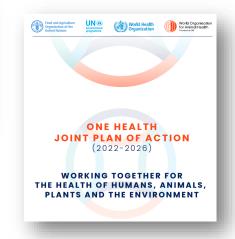






<u>Surveillance and information sharing operational tool:</u> <u>an operational tool of the tripartite zoonoses guide</u> <u>Sept</u>





The Food and Agriculture Organization of the United Nations (FAO)

Emerging zoonotic coronaviruses in animals' situation updates



Emerging Zoonotic Coronaviruses in animals - situation update FAO Animal Health Service / EMPRES



Updates

September 2025

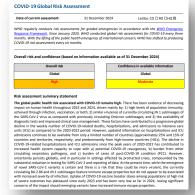
- Two new MERS-CoV human cases reported in Saudi Arabia, with one fatality;
- SARS-CoV-2 virological findings reported in animals in Argentina, Brazil and Bulgaria;
- · MERS-CoV virological findings reported in animals in Kenya;
- 3 new species reported naturally infected by SARS-CoV-2;
- 13 new relevant publications.

Emerging zoonotic coronaviruses in animals



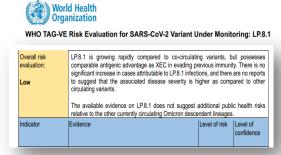
Guidelines, Reports and Risk Assessments

COVID-19 Global Risk Assessment



COVID-19 Global Risk Assessment

SARS-CoV-2 Variant Risk Evaluations



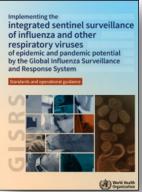
Tracking SARS-CoV-2 variants

COVID-19 Vaccination Insights Report



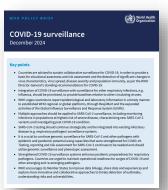
COVID-19 Vaccination Insights
Report

Integrated surveillance



Implementing the integrated sentinel surveillance of influenza and other respiratory viruses of epidemic and pandemic potential by the Global Influenza Surveillance and Response System: standards and operational guidance

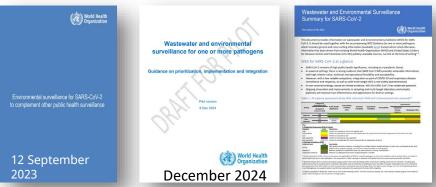
COVID-19 policy briefs



COVID-19 policy briefs

(6 Topics: Surveillance, Testing, Clinical management, Vaccination, IPC, RCCE)

Wastewater Surveillance



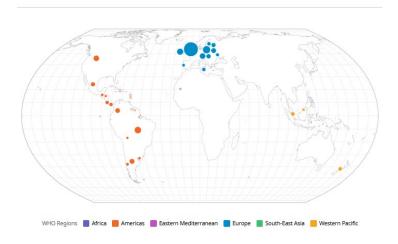
Environmental surveillance for SARS-CoV-2 to complement other public health surveillance Water Sanitation and water Sanitation and water Sanitation and programme



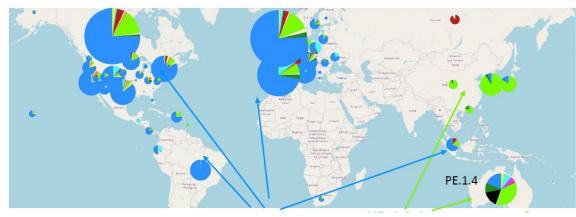
Key areas for strengthening

- Most available data comes from high-income settings.
- Limited geographical and income-level representativeness in assessing burden and variant surveillance.
- Evidence based decision making, guidance development and risk assessments are highly dependent to the information from limited geographical areas and high-income context.
- Countries from underrepresented regions are encouraged to monitor the burden of COVID-19 and SARS-CoV-2 variants and share information with global public health community.

Number of hospitalizations due to SARS-CoV-2 in last 28 days



SARS-CoV-2 variant sequence submissions Later: 2025-09-10 to 2025-10-20 (7045)



Courtesy of Bette Korber

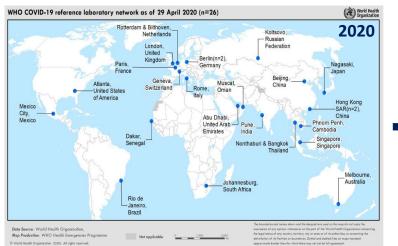




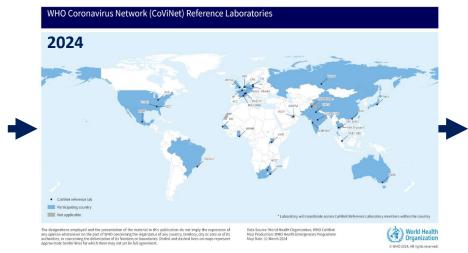


WHO COVID-19 reference laboratory network evolved to the WHO Coronavirus Network





Labs from 26 countries across WHO regions



network is expected to expand to include more than 48 laboratories across 35 countries.

The second expression of interest was

launched in 2025, and the selection process is currently underway. The

34 laboratories 21 countries

- •Human health 27 reference laboratories;
- •Animal health 2 FAO reference labs and 2 WOAH Collaborating Centres = 4
- •Environment -2 WHO CC +1 (consortium) =3

Objectives:

- Facilitate early and accurate detection
- Support surveillance and monitoring of the global circulation and evolution
- Provide timely risk assessment to inform WHO policy related to a range of public health and medical countermeasures
- Support capacity building of laboratories relevant to needs of WHO and CoViNet, particularly those in low and middle-income countries

a organization

Activities 2024-2025

- ■SARS-CoV-2 antigenic characterization work for timely risk assessment facilitated by the WHO Biohub,
- Building advanced diagnostic capacity (neutralisation assays for coronaviruses) in Ghana and Pakistan
- ■Boosting SARS-CoV-2 sequencing in Africa (Uganda/South Sudan and Senegal)
- •Harmonised protocols for panCoV detection by RT-PCR shared with CoViNet laboratories
- External Quality Assessment for zoonotic coronavirus sent to over 30 CoViNet reference laboratories

SARS-CoV-2 Evolution

An early warning system for emerging SARS-CoV-2 variants

Global sequencing and surveillance capacity for SARS-CoV-2 must be strengthened and combined with multidisciplinary studies of infectivity, virulence and immune escape, in order to track the unpredictable evolution of the ongoing COVID-19 pandemic.

Lorenzo Subissi, Anne von Gottberg, Lipi Thukral, Nathalie Worp, Bas B. Oude Munnink, Surabhi Rathore, Laith J. Abu-Raddad, Ximena Aguilera, Erik Alm, Brett N. Archer, Homa Attar Cohen, Amal Barakat, Wendy S. Barclay, Jinal N. Bhiman, Leon Caly, Meera Chand, Mark Chen, Ann Cullinane, Tulio de Oliveira, Christian Drosten, Julian Druce, Paul Effler, Ihab El Masry, Adama Faye, Simani Gaseistiwe, Elodie Ghedin, Rebecca Grant, Bart L Haagmans, Belinda L. Herring, Shilpa S. Iyer, Zyleen Kassamali, Manish Kakkar, Rebecca J. Kondor, Juliana A. Leite, Yee-Sin Leo, Gabriel M. Leung, Marco Marklewitz, Sikhulile Moyo, Jairo Mendez-Rico, Nada M. Melhem, Vincent Munster, Karen Nahapetyan, Djin-Ye Oh, Boris I. Pavlin, Thomas P. Peacock, Malik Peris, Shibin Peng, Leo L. M. Poon, Andrew Rambaut, Jilian Sacks, Vinzbong Shen, Marilda M. Siqueira, Sofonias K. Tessema, Erik M. Volz, Volker Thiel, Sylvie van der Werf, Sylvie Briand, Mark D. Perkins, Maria D. Van Kerkhove, Marion P. G. Koopmans and Anurag Agrawal

Comment

https://doi.org/10.1038/s41591-024-02949-0

An updated framework for SARS-CoV-2 variants reflects the unpredictability of viral evolution

Lorenzo Subissi, James Richard Otieno, Nathalia Worp, Homa Attar Cohen, Bas B. Oude Munnink, Laith J. Abu-Raddad, Erik Alm, Amal Barskat, Wendy S. Barclay, Jinal N. Bhiman, Leon Caly, Meera Chand, Mark Chen, Ann Cullinane, Tulio de Oliveira, Christian Drosten, Julian Druce, Paul Effler, Ihab El Masry, Adama Faye, Elodie Ghedin, Rebecca Grant, Bart L. Haagmans, Christian Happi, Belinda L. Herring, Emma B. Hodoroft, Juniorcaius Ikejezie, Victoria Katawera, Zyleen Alnashir Kassamali, Yee-Sin Leo, Gabrie M. Leung, Rebecca J. Kondor, Marco Marklewitz, Alairo Mendez-Rico, Nada M. Melhem, Vincent Munster, Karen Nahapetik, Jario Mendez-Rico, Nada M. Melhem, Vincent Munster, Karen Nahapetik, Jenio Peng, Leo L. M. Poon, Andrew Rambaut, Senjuti Saha, Vinzhong Shen, Marilda M. Siqueira, Erik Volz, Sofnais K. Tassema, Volker Thiel, Henda Triki, Sylvie van der Werf, Karin von Ejie, Jane Cunningham, Marion P. G. Koopmans, Anne von Gottberg, Anuraa Acrawa & Maria D. Van Kerkhove

Check for updates



- SARS-CoV-2 variant risk evaluation in place since the emergence of variants
- Framework formally published in August 2023, and then updated, under the advice of the Technical Advisory Group on Virus Evolution (TAG-VE)





Technical Advisory Group on Virus Evolution (TAG-VE)

What it is:

An independent, multidisciplinary advisory group assessing public-health implications of priority viruses and informing WHO policy.

Function:

- 1. Determines which variants warrant further investigations
- 2. Assesses whether variants alter
 - -transmission or disease characteristics or
 - -performance of vaccines, therapeutics, diagnostics or
 - -effectiveness of public health and social measures

Outputs (2025):

MERS-CoV risk evaluation (27-10-2025)

Mpox

Clade Ia risk evaluation (24-01-2025) Clade Ib risk evaluation (24-01-2025)

SARS-CoV-2

LP.8.1 (03-02-2025)

NB.1.8.1 (23-05-2025)

XFG (25-06-2025)

Oropouche (10-06-2025)





SARS-CoV-2 Risk Evaluation for XFG

Workflow:

- 1. Surveillance and genomics inputs
- 2. Expert review
- 3. Risk evaluation
- 4. Recommendations/comm unication





VOC, VOI, and VUM definitions and actions (updated 4 Oct 2023)





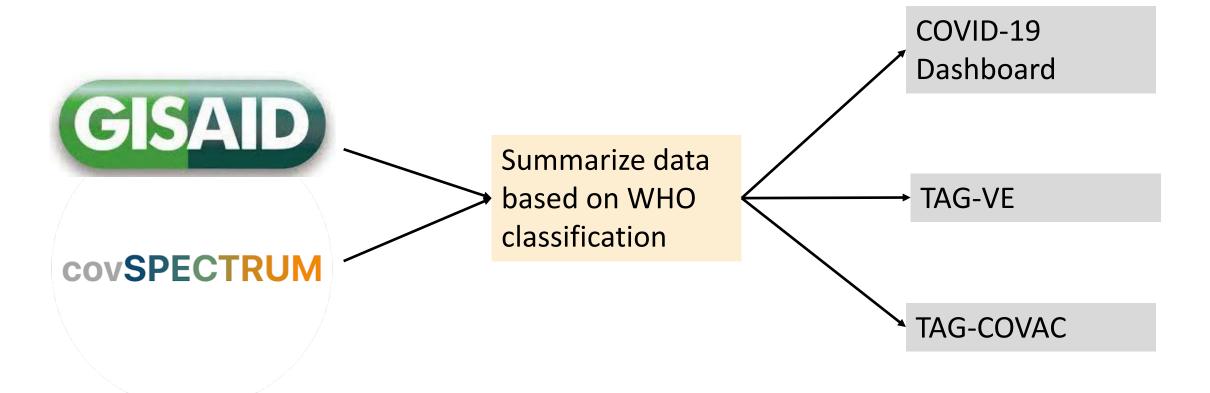
TAG-VE: SARS-CoV-2 Variant Designation and Risk Evaluation, January 2024 to October 2025

 Lack of crucial studies, such as those that were on variant disease severity





Genomic surveillance data source and pipeline







Challenges: Number of Member States sharing genomic sequencing data





Technical Advisory Group on COVID-19 Vaccine Composition (TAG-CO-VAC)



Dr Bart Haagmans, TAG-CO-VAC Chair

EPI-WIN Webinar

COVID-19 vaccines

• Fact: COVID-19 vaccines developed early in the pandemic provide high levels of protection, particularly against severe disease and death.

 Problem: SARS-CoV-2 continues to evolve to escape vaccine induced immunity and vaccine induced immunity wanes in time.

 Solution: COVID-19 vaccines are updated to respond to circulating SARS-CoV-2 variants.







WHO Advisory Groups for COVID-19

Aim: Monitor and assess SARS-CoV-2 variants and evaluate their impact on countermeasures, including vaccines, therapeutics, diagnostics or effectiveness of public health and social measures.



Monitoring & surveillance

TAG-VE

- Determines which variants warrant further investigations
- · Assesses whether variants alter
 - transmission or disease characteristics or
 - performance of vaccines, therapeutics, diagnostics or
 - effectiveness of public health and social measures



Research, evidence & assessment

R&D Blueprint for epidemics

 COVID-19 Vaccines Research Expert Group

TAG-CO-VAC

- Assesses the impact of SARS-CoV-2 variants on current COVID-19 vaccines
- Determines whether changes to vaccine composition are needed





SAGE

 Recommends policies and strategies on vaccine use and immunization programmes through evidence-based assessment







TAG-CO-VAC Members

Chair



Dr Bart Haagmans

Associate professor at the Department of Viroscience of Erasmus MC, Rotterdam, the Netherlands

Learn more >

Vice-Chair



Professor Cheryl Cohen

Professor in epidemiology at the University of the Witwatersrand and Head of the Centre for Respiratory Disease and Meningitis at the National Institute for Communicable Diseases

Learn more >

TAG-CO-VAC Secretariat:

tagcovac@who.int

Members

Dr Svein Rune Andersen >

Scientific Director for Vaccines at the Norwegian Institute of Public Health, Norway



Dr Janine Kimpel >

Group leader and deputy Director at the Institute of Virology at the Medical University of Innsbruck, Austria



Prof Narendra Kumar Arora >

Executive Director of The INCLEN Trust International, India



Professor at the Population Health Group at the National Centre for Immunisation Research and Surveillance (NCIRS) in Australia



Prof Laith Abu-Raddad >

Prof of Infectious Disease Epidemiology at Weill Cornell Medicine-Qatar, Cornell University,



Professor Elizabeth Miller >

Professor in Infectious Disease Epidemiology at the London School of Hygiene and a visiting professor at the Sackler School of Public Health at Tel Aviv University



Lead of the Infection, Epidemiology and Policy Analytics Group at the Kirby Institute, University of New South Wales

Technical Adviser for Capacity Building and Clinical Evaluation of COVID-19 vaccines



Dr Irene Owusu Donkor >

Associate Professor Deborah Cromer >

Fellow in the Epidemiology Department of the Noguchi Memorial Institute for Medical Research (NMIMR)



Professor of Microbiology and Immunology, and of Pediatrics at the University of Iowa



Professor Kanta Subbarao >

Director of the WHO Collaborating Centre for Reference and Research on Influenza and Professor, Department of Microbiology and Immunology



Prof Derek Smith >

Director of the Center for Pathogen Evolution, and Professor of Infectious Disease Informatics at the University of Cambridge in the United Kingdom



Director of the WHO Collaborating Centre for Reference and Research on Influenza, Japan



Director-in-Charge of the National Institute of One Health, Nagpur, and a Scientist 'F' at the Indian Council of Medical Research-National Institute of Virology (ICMR-NIV), Pune, India





https://www.who.int/groups/technical-advisory-group-on-covid-19-vaccine-composition-(tag-co-vac)/about





WHO Technical Advisory Group on COVID-19 Vaccine Composition

Functions of the TAG-CO-VAC:

- Make recommendations to WHO on the methods to assess the impact of SARS-CoV-2 variants on COVID-19 vaccines;
- Provide interpretation of available evidence on the effect of SARS-CoV-2 variants on COVID-19 vaccines, including but not limited to vaccine effectiveness;
- Recommend to WHO, for each COVID-19 vaccine platform, adaptations (if any) needed so that vaccines continue to safely provide protection against SARS-CoV-2 variants.

Meeting frequency:

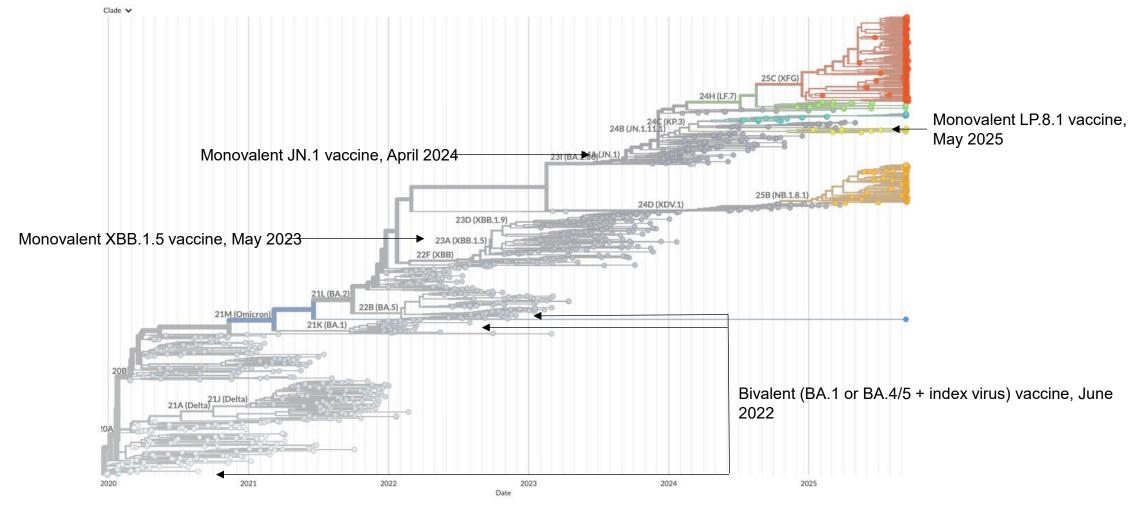
• The TAG-CO-VAC continues to convene closed, decision-making meetings approximately every six months. After each meeting, recommendations to either maintain current vaccine composition or to consider updates are issued.







SARS-CoV-2 evolution and COVID-19 vaccine composition decisions



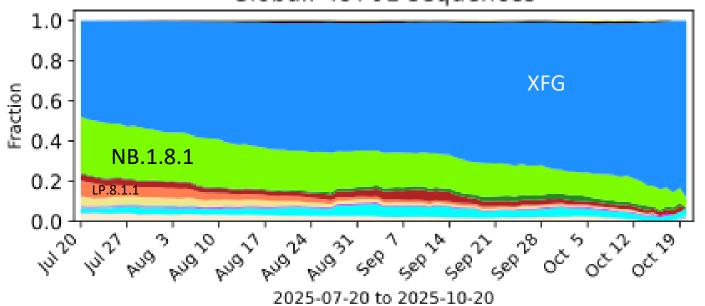






Type of data	Comments
SARS-CoV-2 genetic evolution	With support from Technical Advisory Group on Virus Evolution (TAG-VE)
	Key variants include the list of Variants of Interest (VOI) and Variants Under Monitoring (VUM). This list is maintained on the WHO website: https://www.who.int/activities/tracking-SARS-CoV-2-variants





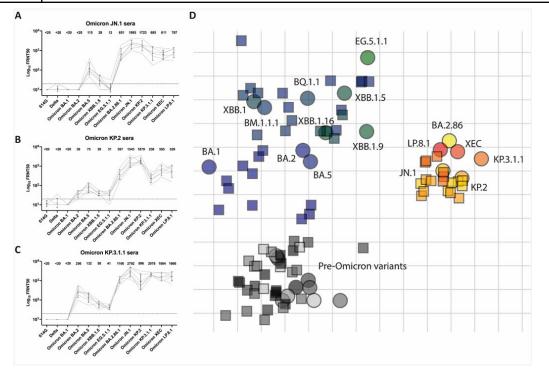
Bette Korber, Los Alamos







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Antigenic characterization of previous and emerging SARS-CoV-2 variants	Animal sera following primary infection or vaccination analyzed in one-way and two-way neutralization tests (pseudotype and live virus neutralization assays).



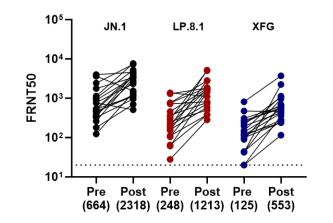






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Antigenic characterization of previous and emerging SARS-CoV-2 variants	Animal sera following primary infection or vaccination analyzed in one-way and two-way neutralization tests (pseudotype and live virus neutralization assays).
Preliminary immunogenicity data on breadth and durability of immune responses following vaccination or infection with SARS-CoV-2 variant antigens	Neutralization of previous and recent variants by non-naïve animal sera (e.g., sequentially immunized or infected);
	Neutralization of previous and recent variants by both pre- and post-vaccination human sera; Neutralization of variants by sera from cohorts that are representative of recent population immunity.





Sentinel - JN.1 vaccination





Type of data	Comments
SARS-CoV-2 genetic evolution	With support from Technical Advisory Group on Virus Evolution (TAG-VE)
	Key variants include the list of Variants of Interest (VOI) and Variants Under Monitoring (VUM). This list is maintained on the WHO website: https://www.who.int/activities/tracking-SARS-CoV-2-variants
Antigenic characterization of previous and emerging SARS-CoV-2 variants	Animal sera following primary infection or vaccination analyzed in one-way and two-way neutralization tests (pseudotype and live virus neutralization assays).
Preliminary immunogenicity data on breadth and durability of immune responses following vaccination or infection with SARS-CoV-2 variant antigens	Neutralization of previous and recent variants by non-naïve animal sera (e.g., sequentially immunized or infected); Neutralization of previous and recent variants by both pre- and post-vaccination human sera; Neutralization of variants by sera from cohorts that are representative of recent population immunity.
Vaccine effectiveness (VE) estimates of currently approved vaccines	Relative VE estimates in human populations of currently approved vaccines. Studies should provide variant-specific estimates and distinct estimates for vaccine antigen compositions across different vaccine platforms.







Types of data considered by TAG-CO-VAC

Type of data	Comments		
SARS-CoV-2 genetic evolution	With support from Technical Advisory Group on Virus Evolution (TAG-VE)		
	Key variants include the list of Variants of Interest (VOI) and Variants Under Monitoring (VUM). This list is maintained on the WHO website: https://www.who.int/activities/tracking-SARS-CoV-2-variants		
Antigenic characterization of previous and emerging SARS-CoV-2 variants	Animal sera following primary infection or vaccination analyzed in one-way and two-way neutralization tests (pseudotype and live virus neutralization assays).		
Preliminary immunogenicity data on breadth and durability of immune	Neutralization of previous and recent variants by non-naïve animal sera (e.g., sequentially immunized or infected);		
responses following vaccination or infection with SARS-CoV-2 variant	Neutralization of previous and recent variants by both pre- and post-vaccination human sera;		
antigens	Neutralization of variants by sera from cohorts that are representative of recent population immunity.		
Vaccine effectiveness (VE) estimates of currently approved vaccines	Relative VE estimates in human populations of currently approved vaccines. Studies should provide variant-specific estimates and distinct estimates for vaccine antigen compositions across different vaccine platforms.		
Data from vaccine manufacturers	Animal and human data that demonstrate the breadth and durability in immune responses elicited by vaccines in current portfolio, as well as any vaccine candidates in development;		
	Observational epidemiological data that demonstrate the efficacy or effectiveness of any vaccines in current portfolio, as well as any vaccine candidates in development.		



TAG-CO-VAC decision-making meeting

Next meeting: December 2025



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Statement on the antigen composition of COVID-19 vaccines

15 May 2025 | Statement | Reading time: 7 min (1811 words)

Key points

- Vaccination remains an important public health countermeasure against COVID-19. As per the WHO
 Director General's standing recommendations for COVID-19, Member States are recommended to continue
 to offer COVID-19 vaccination based on the recommendations of the
 WHO Strategic Advisory Group of
 Experts on Immunization (SAGE).
- SARS-CoV-2 continues to undergo sustained evolution since its emergence in humans, with important genetic and antigenic changes in the spike protein.
- The objective of an update to COVID-19 vaccine antigen composition is to enhance vaccine-induced immune responses to circulating SARS-CoV-2 variants.
- The WHO Technical Advisory Group on COVID-19 Vaccine Composition (TAG-CO-VAC) advises manufacturers
 that monovalent JN.1 or KP.2 vaccines remain appropriate vaccine antigens; monovalent LP.8.1 is a
 suitable alternative vaccine antigen.
- In accordance with WHO SAGE policy, vaccination should not be delayed in anticipation of access to
 vaccines with an updated composition.

Related

Annex: Statement on the antigen composition of COVID-19 vaccines

News



Statement on the antigen composition of COVID-19 vaccines 15 May 2025

Annex: Statement on the antigen composition of COVID-19 vaccines

15 May 2025

Evidence to support considerations of COVID-19 vaccine antigen composition

The data highlighted below are representative examples of the data reviewed and considered by the TAG-CO-VAC to inform the recommendation on COVID-19 vaccine composition and include:

- SARS-CoV-2 genetic evolution;
- (2) Antigenic characterization of previous and emerging SARS-CoV-2 variants using virus neutralization tests with animal antisera and further analysis of antigenic relationships using antigenic cartography;
- (3) Immunogenicity data on the breadth of neutralizing antibody responses elicited by currently approved vaccine antigens against circulating SARS-CoV-2 variants using animal and human sera;
- (4) Preliminary immunogenicity data on immune responses following infection with circulating SARS-CoV-2 variants;
- (5) Available vaccine effectiveness (VE) estimates of currently approved vaccines during periods of JN.1 lineage circulation; and
- (6) Preliminary non-clinical and clinical immunogenicity data on the performance of candidate vaccines with updated antigens shared confidentially by vaccine manufacturers with TAG-CO-VAC (data not shown).

The TAG-CO-VAC convenes a Subgroup comprised of Members and Advisors with virological and immunological expertise. The data highlighted below were also reviewed and considered by the Subgroup. Confidential data reviewed by the TAG-CO-VAC and the Subgroup are not shown.

1. SARS-CoV-2 genetic evolution

SARS-CoV-2 continues to evolve since its emergence in humans, with important genetic and antigenic evolution of the spike protein.

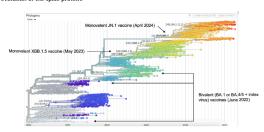


Figure 1: Phylogeny of SARS-CoV-2 variants since its introduction in humans illustrated using Nextsrain. Time is shown on the x axis and selected named variants are labeled with their Nextsrain clade (Pango lineage) at their root nodes. Clades included as vaccine antigens are indicated with the date of the TAG-CO-VAC recommendation for this vaccine antigen composition.







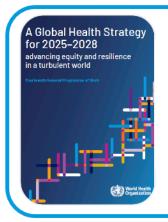
Strategic Advisory Group of Experts (SAGE) on Immunization

Informing decisions on vaccination – the work of SAGE

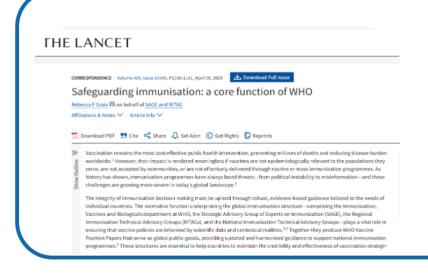
Annelies Wilder-Smith MD PhD MPH, Executive Secretary, SAGE Secretariat



WHO and SAGE's normative role in immunization policy is more critical than ever



- WHO's 2025-28 strategy: 'developing evidence-based norms and standards' a key component for advancing equity and resilience in a turbulent world
- WHO's prioritization exercise highlighted immunization and policy as core function



The normative function underpinning the global immunization structure (SAGE, RITAGS & NITAGS) play a vital role in ensuring vaccine policies are informed by scientific data & contextual realities

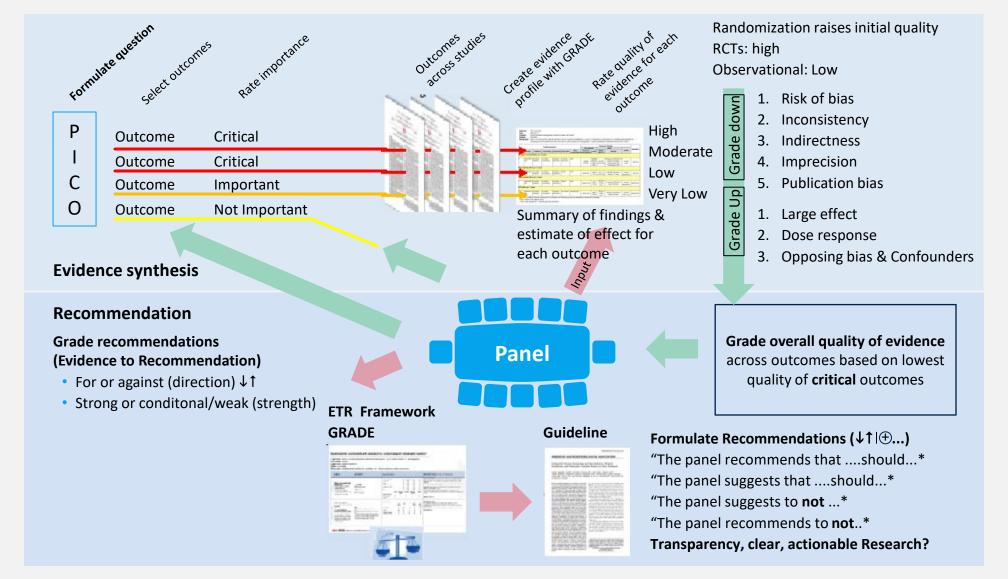
Key Attributes

- Transparency
- Independence
- Best available evidence
- Evidence to policy methodology
- Support country decision-making in evolving contexts



- Identification of problem/ Terms of Reference/ Establishment of Working Group
- Systematic review of the literature
 - A. Definition of critical questions
 - B. Systematic search
 - C. Assessment of risk of bias
 - D. GRADE
- Development of evidence to recommendation table
- 4 Draft recommendations
- 5 Presentation to SAGE
- 6 SAGE discussion, deliberation and decision
- 7 Publication as WHO vaccine position paper

GRADE approach





Evidence to Decision Table Criteria

- Problem Statement
- Benefits and harms plus GRADE
- Values and Preferences
- Resource use and cost-effectiveness
- Equity impacts
- Acceptability
- Feasibility

Concluding with

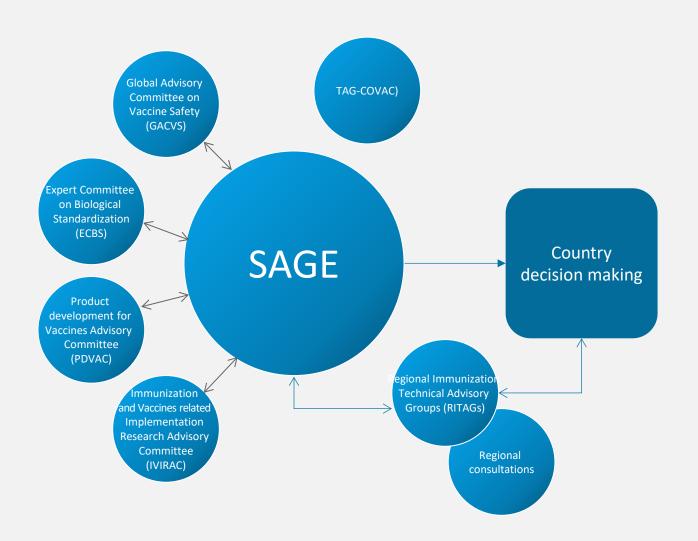
Balance of Benefits and Harms, Type of Recommendation and Recommendation Text







SAGE and other WHO advisory groups



WHO Interim Recommendations for the optimal use of COVID-19 vaccination

HIGH priority-use groups					
Target population	Vaccination of persons who have never received a COVID-19 vaccine	Revaccination of persons who have received at least one dose of COVID-19 vaccine			
Oldest adults ⁱ	Single dose ⁱⁱⁱ	6-12 months after previous dose			
Older adults with multiple comorbidities that put them at higher risk of severe COVID-19					
Older adults ⁱⁱ	Single dose ⁱⁱⁱ	Approximately 12 months after previous dose			
Other adultsiv with severe obesity or a comorbidity that puts them at higher risk of severe COVID-19					
MEDIUM priority-use groups					
Target population	Vaccination of persons who have never received a COVID-19 vaccine	Revaccination of persons who have received at least one dose of COVID-19 vaccine			
Healthy adults ^{iv}	Single dose ⁱⁱⁱ	Not routinely recommended ^{vi}			
Children and adolescents aged 6 months to 17 years with severe obesity or a comorbidity that puts them at higher risk of severe COVID-19 ^v					

LOW priority-use groups				
Target population	Vaccination of persons who have never received a COVID-19 vaccine	Revaccination of persons who have received at least one dose of COVID-19 vaccine		
Healthy children and adolescents aged 6 months to 17 years	If countries opt to vaccinate low priority-use groups ^{vii} , they could consider single dose for ages 5 years and above; two doses for age 6 months to 4 years ^v	Not routinely recommended ^{vi}		

Sub-populations with special considerations					
Target population	Vaccination of persons who have never received a COVID-19 vaccine	Revaccination of persons who have received at least one dose of COVID-19 vaccine			
Persons with moderate and severe immunocompromising conditions (adults, adolescents and children > 6 months)	Two or three doses in consultation with the health care provider	6-12 months after previous dose; optimal time interval should be determined in consultation with the health care provider			
Pregnant adults and pregnant adolescents ^{viii}	Single dose in each pregnancy regardless of previous vaccination status; ideally during the second trimester or at any opportunity				
Health and care workers with direct patient contact	Single dose	Approximately 12 months after previous dose			

i Age cut-off to be decided by countries; often it is 75 or 80 years. ii Age cut-off to be decided by countries: often it is 50 or 60 years. iii In vaccine-naïve persons, for programmatic purposes, a single dose can be considered for primary vaccination given that the vast majority of the population will have been infected at least once. For inactivated COVID-19 vaccines, two doses are required for the primary vaccine series. iv Age cut-off to be decided by countries: often it is 18 to 49 or 18 to 59 years. v Regulatory approvals or WHO EUL for the age indication differ by vaccine product; refer to the product-specific vaccine recommendations. vi "Not routinely recommended" means that such vaccines are not recommended because of minimal public health impact and low cost-effectiveness in most settings. However, vaccination may be offered in individual country-specific circumstances where added benefit is expected to be more substantial. This interim recommendation acknowledges that some countries may elect to offer such doses in the routine programme based on population risks, disease epidemiology, or public health priorities. vii Benefit of vaccinating healthy children and adolescents is substantially lower compared to vaccinating older persons or as compared to other childhood vaccinations. Countries could consider vaccination based on disease burden, cost effectiveness, and other programmatic priorities. viii Regulatory approvals or WHO EUL for the use in pregnancy may differ by vaccine product.

Proposed policy questions in the era of high population level immunity and Omicron sublineages

Should WHO continue to recommend 12 monthly boosters to high-priority use groups?

Should the most vulnerable populations (e.g., the very elderly and those with multiple comorbidities) continue to receive boosters every 6–12 months, or could this be adjusted to a 12-month intervals?

Should WHO continue to recommend 12 monthly boosters to health workers regardless of age and comorbidities?

Should WHO continue to recommend COVID-19 vaccination for each pregnancy?

Should WHO continue to not recommend annual boosters for medium- and low-priority use groups?



Systematic review of literature



PICO question: re-vaccination with a variant-adapted vaccine in past year versus no vaccination in past year



Assessment of the risk of bias of retrieved evidence



Rating of the quality of the evidence (GRADE)



Global status of COVID-19 vaccination programmes and resources available to support programme strengthening

Shoshanna Goldin – Technical Officer

5 November 2025

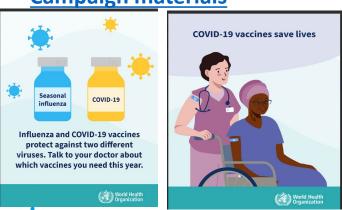


COVID-19 Vaccination Data (2024) and Tools

Target population group	No. of Member States with a policy for COVID-19 revaccination/boosters	No. of Member States reporting COVID-19 vaccination coverage data	Median coverage of reporting Member States
Older adults	108	42	6.5%
People with chronic conditions	102	17	1%
Health workers	90	20	7.2%
Pregnant persons	81	16	2.7%
Total COVID-19 vaccination (not	58 recommend for all adults	31	3.94%
disaggregated by target	36 recommend for		
population group)	children/adolescents		

Campaign materials

World Health Organization



Coverage manual



Toolkit - guidance, training, and tools





How to achieve high uptake of COVID-19 vaccines?

1. Design data-driven and evidence-based interventions

- Support systematic collection and use of data on behavioural and social drivers
- Monitor and evaluate to guide continuous improvement

2. Target the structural and service-level barriers

- Improve <u>service quality</u>, <u>ease of access</u>, <u>and reduce costs</u>, e.g., meet their needs of priority populations
- Integrate vaccination in primary care and relevant adult vaccination platforms, especially in disadvantaged settings

3. Engage meaningfully with communities and civil society

- Genuinely <u>listen</u>: create spaces for dialogue to explore and address concerns
- Equip and empower advocates to promote positive norms and demonstrate the importance of vaccination
- Co-design interventions to increase outcomes and local ownership, e.g., human-centred design

4. Tailor communications and work through trusted messengers

- Communicate in ways that are honest, transparent, timely, credible, relatable and engaging
- Be ready to respond to vaccine-related events, e.g., AEFI, anti-vaccination activism

Tools and guidance: <a href="https://www.who.int/initiatives/act-accelerator/covax/covid-19-vaccine-country-readiness-and-delivery/acceptance-and-demand-demand-demand-delivery/acceptance-and-demand-demand-demand-delivery/acceptance-and-demand-demand-demand-delivery/acceptance-and-demand-demand-delivery/acceptance-and-demand-delivery/acceptance-and-demand-delivery/acceptance-and-delivery/acceptance-and-delivery/acceptance-and-demand-delivery/acceptance-and-de



