



# Monitoring and evaluating public health pathogen genomics

South and Southeast Asia Pathogen Genomics Prioritization and Implementation Workshop

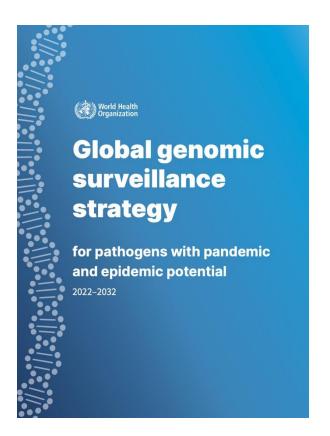
12 September 2024

## Monitoring and evaluating pathogen genomics





Monitoring and evaluation is key to understand progress towards and drive the achievement of the strategy's results



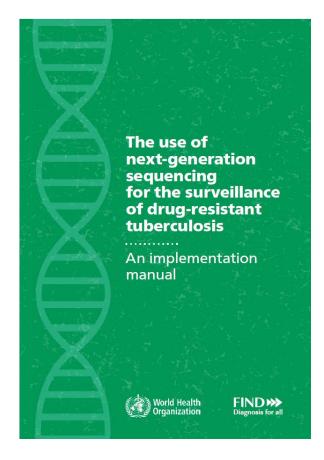
Global genomic surveillance strategy for pathogens with pandemic and epidemic potential, 2022–2032

## Monitoring and evaluating pathogen genomics





A framework for M&E of the impact of NGS is essential to inform decision-making.



The use of next-generation sequencing for the surveillance of drugresistant tuberculosis:

an implementation manual. Geneva: World Health Organization; 2023

## Purpose of evaluating public health pathogen genomics





#### **Inefficiencies and Redundancies**

Identify and reduce inefficiencies or redundancies in the system



#### **Guidelines and Policies**

Development of appropriate guidelines and policies around the use of genomics in public health practice

#### **Equal Distribution**

Ensure public health benefits are equitably distributed



Purpose



#### Sequencing Capacity

Development of sequencing capacity in a considered, informed way

#### **Unexpected Outcomes**

Identify unexpected outcomes (either positive or negative) and take appropriate action as necessary



<u>-\$-</u>1

### Investment, Infrastructure and Training

Identify needed investment, infrastructure and training



### **Expected Outcomes of Implementation**

Identify unexpected outcomes (either positive or negative) and take appropriate action as necessary

## Monitoring and evaluating pathogen genomics



- Guidance on M&E is lacking
- Strong focus on sequencing capacity, rather than public health implementation

For cross-cutting aspects of this strategy, a number of high-level measures should also be monitored to ensure that all countries have access to genomic surveillance. Measures including the following:

- Countries with in-country capability to perform next generation sequencing.
- Countries sharing genomic data to publicly accessible databases or as guided by WHO programmes.
- Countries participating in global quality assessment programmes for sequencing and bioinformatics.
- Countries participating in surge exercises to test genomic surveillance systems.

Global genomic surveillance strategy for pathogens with pandemic and epidemic potential, 2022–2032

### Pathogen Genomics in Public Health Surveillance Evaluation Framework



Phase 2: Reporting and

communication

Analysis and reporting processes

Utility of information presented to

Effectiveness of reporting formats

in information retention

Sequence data sharing and

Sequence data portability and



Phase 3: Public health implementation

#### Part 1

- Perceived utility and utilisation of WGS in public health practice
- Level of articulation between WGS and public health activities
- Facilitators and barriers to integration of sequence data in public health practice
- Use of WGS in public health decision-making

#### Part 2

- Impacts of WGS on public health outcomes
  - Number of notifiable cases
  - Mortality due to notifiable illnesses
  - Cases averted
  - Size of clusters
  - Size of cluster:
  - Spread of clusters
  - Percent of cases linked to clusters
  - Health care costs
  - Costs of epidemiological investigation
  - Loss due to food recalls

#### Data collection:

- Interviews with end users
- Case studies
- Routinely collected public health data
- Variable according to pathogen (see Supplementary materials)

- 'Whole of system' approach
- Focus on implementation in public health
- Three pathogen case studies:
  - Listeria monocytogenes
  - Mycobacterium tuberculosis
  - SARS-CoV-2

· Sample collection and preparation

Phase 1: Pre-analysis and analysis

- Changes to laboratory workflow
- Costings
- Efficiency
- Turnaround times
- Quality of information provided by WGS

#### Data collection:

- Routinely collected laboratory data
- Interviews with laboratory personnel
- Interviews with end users

## Ferdinand AS, Kelaher M, Lane CR, da Silva AG, Sherry NL, Ballard SA, Andersson P, Hoang T, Denholm JT, Easton M, Howden BP, Williamson DA. An implementation science approach to evaluating pathogen whole genome sequencing in public health. Genome Med. 2021 Jul 28;13(1):121. doi: 10.1186/s13073-021-00934-7. PMID: 34321076: PMCID: PMC8317677.

#### Data collection:

in place

end users

governance

interoperability

- · Interviews with bioinformaticians
- Interviews with end users
- Developed reports

### What we've learned: The literature



- Limited literature on evaluation of pathogen genomics
  - PG-PHASE is the only available pathogen genomics evaluation framework
  - Economic evaluations are small-scale, limited in scope and frequently not full cost-benefit analyses
  - No evaluations of the functionality or processes of pathogen genomicsinformed surveillance

Ferdinand AS, Kelaher M, Lane CR, da Silva AG, Sherry NL, Ballard SA, Andersson P, Hoang T, Denholm JT, Easton M, Howden BP, Williamson DA. An implementation science approach to evaluating pathogen whole genome sequencing in public health. Genome Med. 2021 Jul 28;13(1):121. doi: 10.1186/s13073-021-00934-7. PMID: 34321076; PMCID: PMC8317677.

Tran, M., Smurthwaite, K. S., Nghiem, S., Cribb, D. M., Zahedi, A., Ferdinand, A. S., Andersson, P., Kirk, M. D., Glass, K., & Lancsar, E. (2023). Economic evaluations of whole-genome sequencing for pathogen identification in public health surveillance and health-care-associated infections: a systematic review. Lancet Microbe, 4(11), e953-e962. https://doi.org/10.1016/s2666-5247(23)00180-5

### What we've learned: Our experience

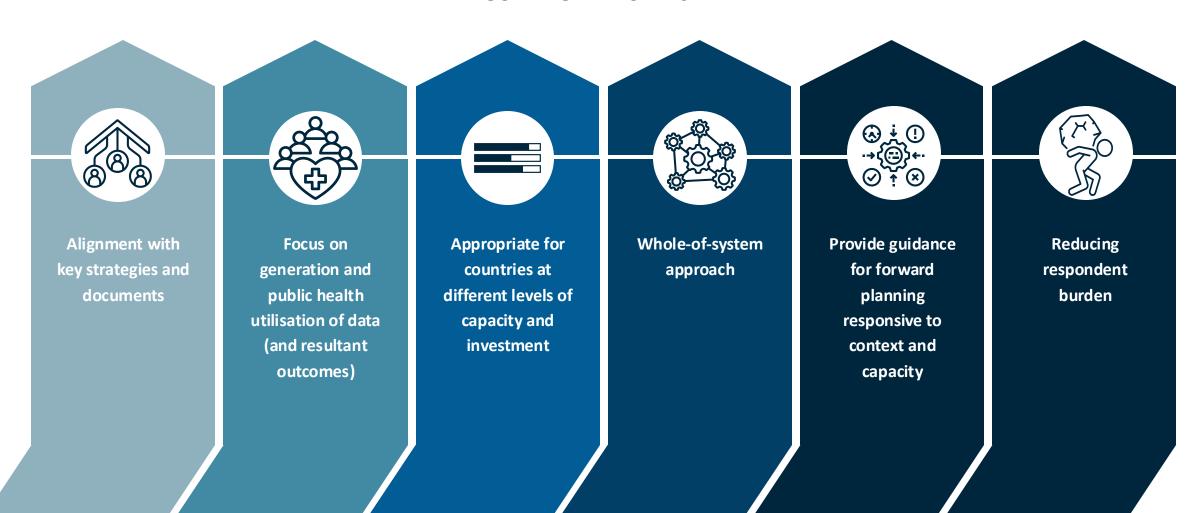


- Need to address respondent burden
- Systems frequently not set up to easily capture relevant data
- Strong focus on sequencing and wet lab capacity
- Limited visibility across different parts of the system (and from different actors)
- Need for tools to clearly signpost areas for strengthening and capacitybuilding

## Developing a pathogen genomics monitoring and evaluation tool



#### **GUIDING PRINCIPLES**



### The 'maturity model' approach



#### **Planning**

- Public health pathogen genomics practices in the planning stages.
- Little to no capacity for or access to sequencing, or sequencing delivers little or no public health benefit.

#### Developing

- Growing but inconsistent public health pathogen genomics practices.
- Sequencing and implementation capacity is being developed, but it is uneven across the system.
- Examples of good practice, but full benefit from sequencing is not realised.

#### **Implementing**

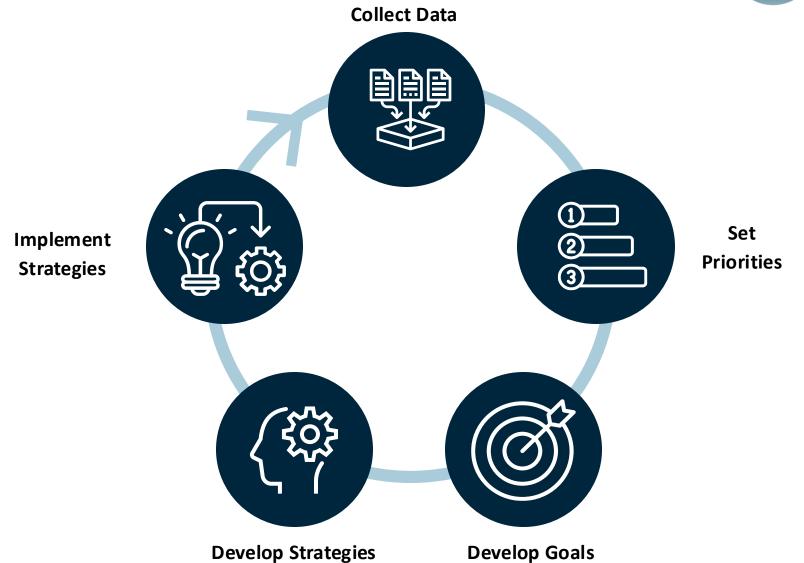
- Practices are established and consistent.
- Pathogen genomics is routinely and appropriately used to inform public health decisionmaking and implementation.
- Benefits are equitably distributed across the population.

### Delivering

- Public health decision-making and policy development strongly benefits from the integration of pathogen genomics data in surveillance.
- The pathogen genomics-informed surveillance system is regarded as an outstanding example.

### The 'maturity model' approach





### **Tool structure**



**Six Sections** 

Genomics-informed surveillance and policy

Specimen selection, collection and referral

**Laboratory workflow** 

**Bioinformatics and analysis** 

Reporting and communication

Implementation in public health practice

Up to six sub sections per section

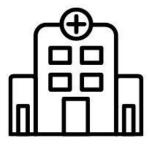
Responding to items in each sub-section provides indication of current maturity level

### **Tool structure**



- For use against a single 'system'
  - May be national for small or centralised countries
  - Sub-national for large or federated countries
- Sections can be completed separately or together
- Some sections to ideally be completed jointly between a public health laboratory and health department







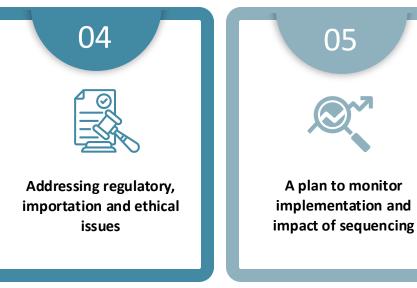
## Section 1: Genomics-informed surveillance and policy











## Section 1: Genomics-informed surveillance and policy



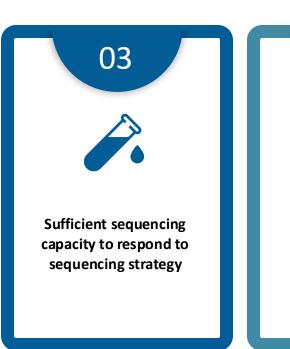
	Planning	Developing	Implementing	Delivering
Sub-section 1.1: Alignment with public health objectives and priorities	Policy to support public health pathogen genomics has not been developed, or there is no alignment between genomics policy and broader public health objectives and priorities	Policies regarding public health pathogen genomics have been developed but are limited in scope and do not align with broader public health objectives and policies	Public health pathogen genomics policy has been developed and there is some socialisation with broader health policy, but alignment is inconsistent	Public health pathogen genomics policy has been developed and socialised to align with identified public health objectives and priorities
Sub-section 1.2: Development of a costed implementation plan	No implementation plan developed	Implementation plan has been developed but no costing data applied	Implementation plan has been developed and costing data applied but with limited transparency	Fully costed implementation plan has been developed with transparent methods of identifying costs
Sub-section 1.3: Required financial resources for implementation and ongoing costs	There is insufficient funding allocated to support key pathogen genomics surveillance (PGS) processes for 1 year	Sufficient funding is secured to support key PGS processes for 1 year	Sufficient funding is secured to support key PGS processes for 2-4 years	Sufficient and sustainable funding for PGS is secured to support key processes for 5+ years
Sub-section 1.4: Addressing regulatory, importation and ethical issues	Policy to support public health pathogen genomics has not been developed, or does not address regulatory, importation or ethical issues	Policies regarding public health pathogen genomics have been developed but are limited in scope and do not address regulatory, importation or ethical issues	Public health pathogen genomics policy has been developed and addresses some regulatory, importation or ethical issues	Public health pathogen genomics policy has been developed and addresses regulatory, importation or ethical issues
Sub-section 1.5: A plan to monitor implementation and impact of sequencing	No monitoring and evaluation plan developed	Monitoring and evaluation plan has been developed but does not take a whole-of-system approach and/or no resources have been allocated	Monitoring and evaluation plan has been developed but does not take a whole-of-system approach or inadequate resources have been allocated	Monitoring and evaluation plan has been developed using a whole-of-system approach and adequate resources have been allocated

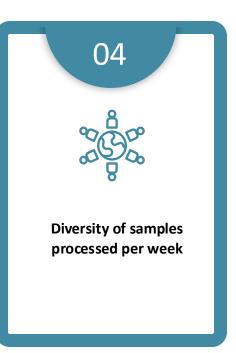
### Section 2: Specimen selection, collection and referral











### **Section 3: Laboratory workflow**





01



Training, competency and retention of laboratory staff

02



Sampling, sample processing capacity and turnaround time

03



Staff, equipment and reagent costs

04



**Quality outcomes** 

05



Participation in global assessment programs for sequencing and bioinformatics

## Section 4: Bioinformatics and analysis



Capacity to use bioinformatics tools

and software to

analyse data

O2

Adequate data storage capacities

Computational power for analytical throughput

Connectivity and ability to connect to international data repositories

Sequence quality assessment processes

Analyse/interpret bioinformatics results

### Section 5: Reporting and communication



01



Interaction with publicly available genomics repositories

02



Staff expertise in genomic epidemiology across laboratory and public health settings

03



Competency and knowledge requirements of end users

04



Appropriateness of information received by end users (i.e., relevance to surveillance objectives, appropriate to enduser level of understanding)

05



System
interoperability to
facilitate data
archiving, tracing
and sharing

06



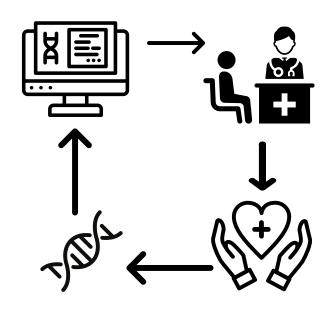
Working relationships between public health laboratories and public health departments

## Section 6: Implementation in public health practice



### Section 6a: Contribution to public health implementation

- Contribution to public health policies and identified priorities
- More precise allocation of investigative resources
- Contribution to appropriately tailored and targeted public health interventions
- Perceptions of affected communities

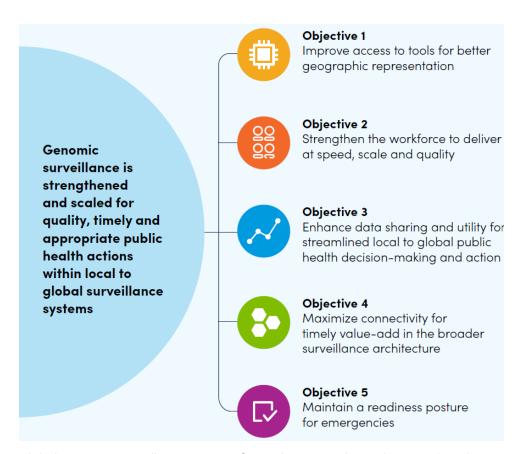


### Section 6b: Quantitative impacts on public health outcomes

- Time between identification of clusters and public health action
- Proportion of cases linked to identified clusters
- Number of notifiable illnesses
- Health care costs due to notifiable illnesses

### Alignment with key guidance documents

### Global genomic surveillance strategy



Global genomic surveillance strategy for pathogens with pandemic and epidemic potential, 2022–2032



### **IPSN Country Capacity Framework**

#### **Hybrid Domestic Capacity**

Leverages partner sequencing and surveillance resources to use genomics for public health decision-making, with limited to no domestic sequencing for monitoring priority pathogens

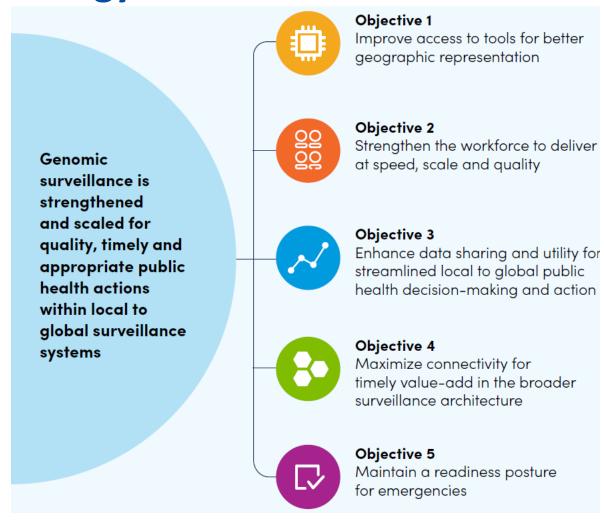
#### **Integrated Domestic Capacity**

Leverages partner resources for strategic support, R&D, supply chain consolidation, and QA, to supplement full domestic sequencing for monitoring priority pathogens

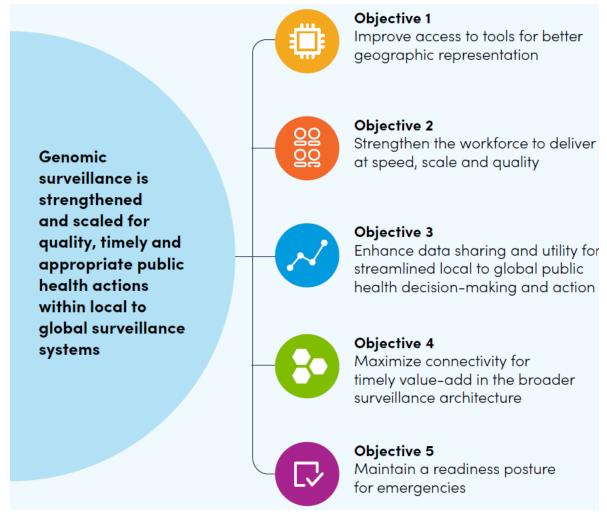
#### Regional/Global Support Capacity

Coordinates regional/global surveillance efforts; provides strategic leadership and support for training, R&D, QA, sequencing, and rapid scale-up; and funds donor programs to enable partner priorities





Global genomic surveillance strategy for pathogens with pandemic and epidemic potential, 2022–2032



Global genomic surveillance strategy for pathogens with pandemic and epidemic potential, 2022–2032

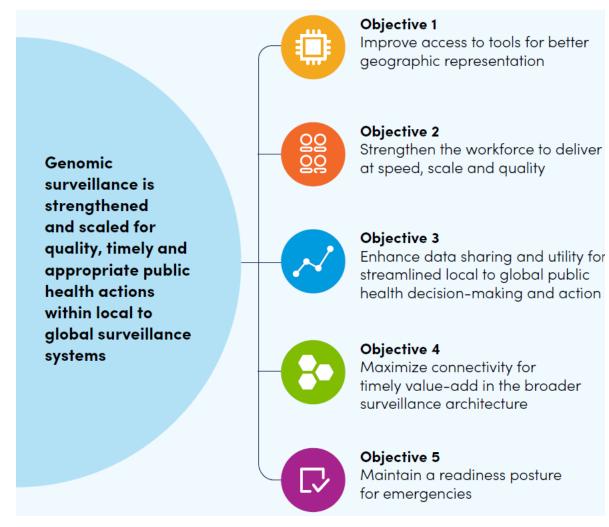




Advocate for and raise awareness of the value of genomic surveillance with policy makers to bring genomics into national disease control strategies.

**Map and monitor capability and capacity landscape** to maximize efficiencies, availability and geographic representativeness.

Define required tools and solutions in order to deliver contextualized, decentralized and sustainable technology and innovation solutions to ensure simple and optimized workflows to enhance access and information sharing.



Global genomic surveillance strategy for pathogens with pandemic and epidemic potential, 2022–2032



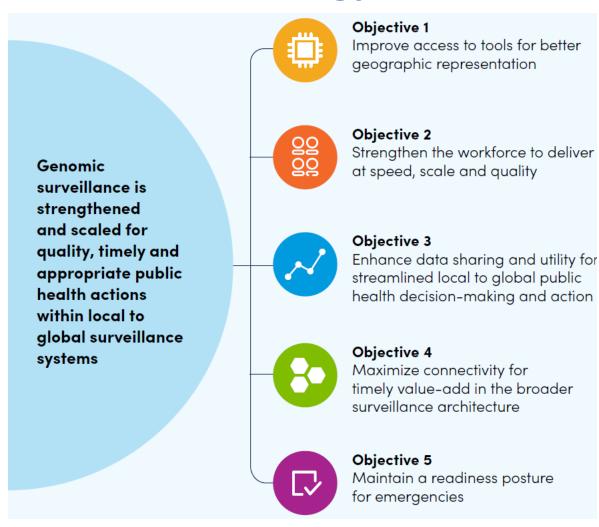




**Develop and roll-out training packages in genomics and bioinformatics** for improved competencies and to facilitate evidencedriven decision making.

Implement external quality assessment programmes for genomics and analytics and provide support to comprehensive quality management systems to ensure accuracy of data and trust in the system.









Develop consensus on **data and meta data standards**, which recognize the importance of data privacy and national sovereignty, while balancing the importance of contextual information to accompany genomic sequencing data.



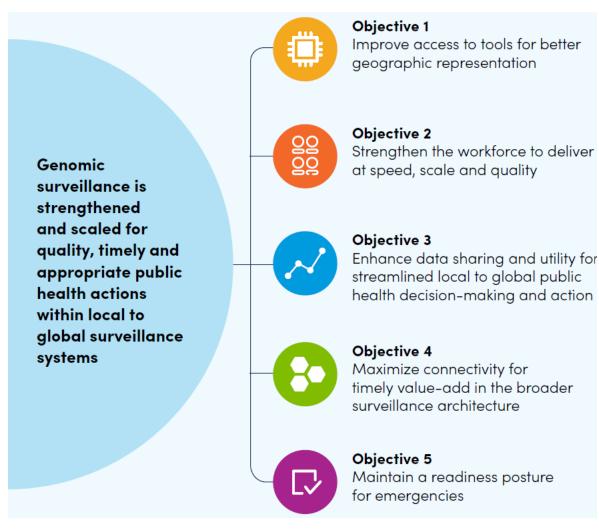
**Establish data sharing and access principles** that are widely agreed upon and explicit to foster transparency for rapid and equitable dissemination.

Ensure **data sharing agreements** are already in place in advance of acute events to promote timely collaboration and coordination.

Harmonize norms, standards, benchmarks, and reference materials to facilitate high quality information sharing.

**Make the use of genomics routine** in surveillance practice and disease prevention, preparedness, readiness and response.











Leverage existing networks to support and facilitate data, specimen and information sharing to foster effective, rapid collaboration to drive public health action.



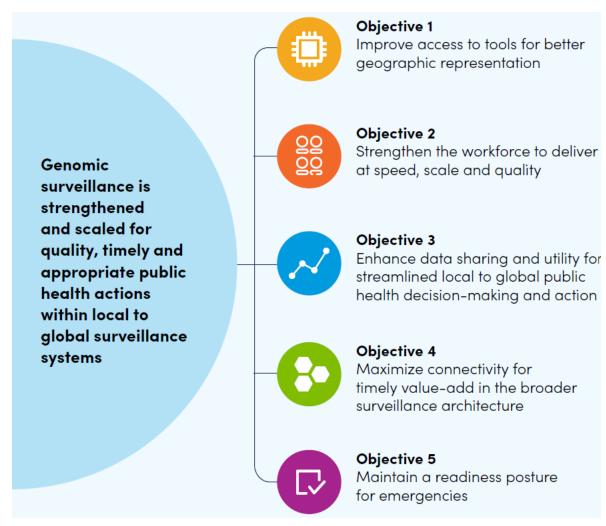
**Increase network linkages** at country, regional and global levels to minimize information siloes and maximize impact, through sharing of resources, protocols and bioinformatics tools.

Implement **targeted collaboration with One Health partners** for comprehensive, integrated surveillance.

Support and strengthen national, regional and global **networks** in routine, epidemic and pandemic contexts.

Global genomic surveillance strategy for pathogens with pandemic and epidemic potential, 2022–2032



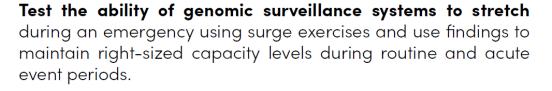














**Establish or sustain joint projects to maintain capacities** and prime systems including the onboarding of new technologies and tools needed at the time of an emergency.

Implement continuous improvement processes including **interor after- action reviews** and utilize information in real-time to strengthen practices.

Global genomic surveillance strategy for pathogens with pandemic and epidemic potential, 2022–2032

## Alignment with IPSN Country Capacity Framework



#### **Hybrid Domestic Capacity**

Leverages partner sequencing and surveillance resources to use genomics for public health decision-making, with limited to no domestic sequencing for monitoring priority pathogens

#### **Integrated Domestic Capacity**

Leverages partner resources for strategic support, R&D, supply chain consolidation, and QA, to supplement full domestic sequencing for monitoring priority pathogens

#### **Regional/Global Support Capacity**

Coordinates regional/global surveillance efforts; provides strategic leadership and support for training, R&D, QA, sequencing, and rapid scale-up; and funds donor programs to enable partner priorities



Assesses domestic capacity to use genomics for public health decision-making



Assesses additional staff capacity to support sequencing strategy and peak sequencing times



Assesses additional instrumentation and reagents to support sequencing strategy and peak periods



Assesses current and potential capacity to support other laboratories as part of a referral system



Assesses overflow capacity in addition to routine genomics work

### **Progress to date and next steps**



Literature review	Completed	
Develop tool structure	Completed	
Develop sub-section maturity model matrices	Completed	
Develop sub-section items	Completed	
Complete tool first draft	Completed	
Consultation, piloting and revision	October 2024	
Finalisation of monitoring and development tool	November 2024	
Further development of documentation, user manual and technical guides; approach for ongoing update as required	January 2025-June 2025	

### **Acknowledgements**



### CPG Pathogen Genomics M&E Tool Working Group

Patiyan Andersson

Mathilda Wilmot

Torsten Seemann

Tilda Thomson

Kristy Horan

Courtney Lane

Hasini Walpola

Sarah Baines

Jake Lacey

Lisa Ionannidis

**Tuyet Hoang** 

Chantel Lin

James Ong

Jessica Ramsey

Jess Gu

Additional SME support provided by

Susan Ballard

### IPSN Monitoring and Evaluation Tool Technical Working Group

Josefina Campos

Tim Dallman

Chantel Lin

Anne E. Purfield

Tuyet Hoang

Noah Hull

Torsten Seemann

Silvia Argimon

Sabrina Weiss Babak Afrough Toni Whistler

Ben Howden

Oluwatosin Tolulope Akane

Swapna Uplekar