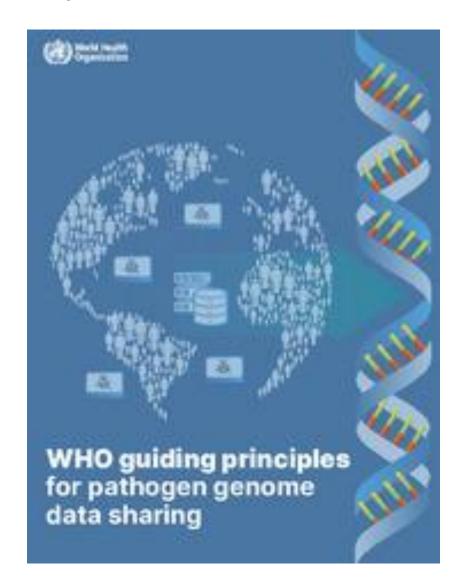
WHO Guiding Principles for Genomic Data Sharing

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- 1. Pathogen genomic initiatives should contribute to capacity development that can establish and sustain data generation, processing, and submission, as well as reliable curation and annotation. In this manner, all countries will be able to analyze data.
- 2. Promote Collaboration and cooperation: between the laboratories and institutions that submit data to it and the scientists who analyze the data and report their analyses
- 3. **Quality** is central In emergencies, there is a trade-off between the time needed to attain very high-quality data and the ability to share data rapidly. In some cases, it may be appropriate to share lower quality data (clearly marked as "preliminary/not fully quality controlled" or similar), with the understanding that preliminary data need to be easily identifiable. The development, implementation, and maintenance of data standards should be prioritized for robust data sharing.
- 4. **Global and regional representativeness**: At the global level, the first few sequences submitted by countries that have not previously shared can be extremely informative. Timeliness and geographical and temporal representation relative to disease epidemiology govern the public health value of pathogen genomic data.



- 5. **Timeliness**: During epidemics and pandemics, ... sharing of ... data should be as timely as possible in order to make data available for analyses. The sharing of pathogen genome data should not be delayed because a data submitter wishes to prepare a scientific manuscript for publication. But neither should data submitters be obliged to waive their rights to the data they share before publication.
- 6. **Acknowledgement and intellectual credit**: All contributions, including those of submitting laboratories, and where appropriate, laboratories from which clinical samples or pathogen isolates have originated, should be appropriately acknowledged in presentations and publications.
- 7. Equitable access to health technologies as a benefit: Data-sharing policies and practices should contribute to equitable access to health technologies wherever possible. Although pathogen genome sequence sharing platforms may not be able to directly ensure access to medical interventions, such platforms should set expectations that all users of the data apply equity and fairness considerations in their use of the data for developing health technologies.



8. As open as possible and as closed as necessary:

Unrestricted open access models for sharing should be used whenever data submitters do not wish rights to be reserved.

Where submitters opt to retain certain protections related to pathogen genome data, platforms that preserve data generators' rights should be used.



9. Interoperability and relevance for decisionmakers:

The global system of pathogen genome data sharing should allow for large-scale standardized analyses that follow good practices developed in a rigorous manner by the global scientific community.

This will permit answers to critical public health questions such as epidemiological characteristics, transmission dynamics, emergence of variants, drug or vaccine resistance and the effectiveness of diagnostics.



- 10. *Trustworthiness and ease of use*: Establishing who controls data sharing and where data are stored is important for achieving trust.
- 11. **Transparency**: Terms and conditions of databases in the global system of pathogen genome data sharing should be publicly available.
- 12. **Consistency with applicable law and ethical regulations**: Pathogen genome data sharing platforms should be established and operated in a manner consistent with applicable national and international laws, regulations, rules, and standards, and ethical regulations, norms, and standards
- 13. **Compliance and enforcement**: Where data sharing mechanisms permit, there may also be a need to raise awareness of instances and sanctions for breaches of the rules of data access as a means of supporting trustworthiness.



Common data sharing questions...

- 1. Will there be adverse repercussions to my sharing the data more openly?
- 2. Will I be scooped if I submit my sequence data to an international platform?
- 3. Will I be acknowledged?
- 4. Will my country have access to medical products developed based on sequences?
- 5. How can sequence sharing align with the open science movement?

Discussion