Network Survey for Safe & Secure Biotechnology Platforms Frequently Asked Questions (FAQs)

Updated as of April 24, 2024

How will potential data be used?

The potential data is primarily intended for use by **researchers** from both the **biotech industry** and **academia**, particularly those focusing on vaccine development or developing AI/ML tools for vaccine design.

Who will have access to the potential data?

Access to the potential data is anticipated to be granted to authorized users, including **researchers**, **healthcare providers**, **device manufacturers** and **software developers** and other appropriate **stakeholders**. The Network Survey seeks to gather input on how to ensure access privileges are appropriately determined, and how user roles, permissions, and compliance are maintained while still allowing for an easy-to-use, highly accessible site.

What are the rules for sharing potential data in the ARPA-H APECx program, including who can access it and where it will be stored?

Researchers selected from APECx must agree to **openly share the potential data acquired during the program**. Any member of the scientific community should have easy access to the potential data, possibly through registration on a repository website. The specific repository will be chosen with input from the ARPA-H Program Manager.

What is the purpose of the Safe & Secure Biotechnology Platforms Network Survey?

The Network Survey for Safe & Secure Biotechnology Platforms aims to solicit feedback from relevant stakeholders to understand the

necessary requirements for end-user adoption. The goal is to hear input directly from potential users to **ensure that the potential APECx toolkit and viral protein data** can be safely and effectively used by the scientific community.

Who should complete the survey?

The program is seeking input from the scientific community, specifically (1) **potential users of the APECx toolkit and datasets**, and (2) **experts in data management, interoperability, storage, and security**. Respondents may come from academic, government, non-profit, startup, or commercial sectors, and may answer as individuals or on behalf of their organization. Relevant areas of experience/expertise include:

- Protein engineering
- Functional and structural protein characterization, multi-omics data
- Data management, applications, interoperability, for large datasets
- Developing and architecting platforms to host data
- Data security and management
- Data harmonization, metadata management, and ontology development
- Vaccine and Ag development
- AI and Machine Learning

Will my responses be kept confidential?

This survey is being administered by the Customer Experience Hub, Advanced Technology International (ATI), part of the Advanced Research Projects Agency for Health (ARPA-H) nationwide health innovation network, ARPANET-H. Contact information provided to complete the survey is confidential and will only be used for survey follow-up. Individual responses to survey questions are confidential and

will only be accessible to select personnel in ARPA-H, and ARPA-H's Customer Experience Hub, ATI. Please do not include medical imaging data, patient data, or personally identifiable information of others in your response. Survey findings and key insights will only be presented in an aggregated format with no identifying information. Findings will be made available to all respondents through a public findings document. This survey does not constitute an endorsement of any initiative from ARPA-H or ATI.

When are responses due?

May 17, 2024.

What is the link? How do I provide input to the survey?

Responses can be submitted here or feel free to scan the QR Code.

When will the public report be published?

We expect to release the public report in July 2024. This is subject to change.