

# Welcome to the National Cybersecurity Center of Excellence

## Virtual Workshop on Exploring Solutions for the Cybersecurity of Genomic Data

Wednesday, May 18, 2022, 1:00 PM – 3:30 PM (EDT)

**We will begin shortly.  
This meeting will be recorded.**

# Virtual Workshop on Exploring Solutions for the Cybersecurity of Genomic Data

Ron Pulivarti  
NIST NCCoE



# Virtual Workshop on Exploring Solutions for the Cybersecurity of Genomic Data

Eric Lin

Director, Material Measurement Laboratory, NIST

# Virtual Workshop on Exploring Solutions for the Cybersecurity of Genomic Data

Robel Worku  
Montgomery County Economic Development  
Corporation



# Priming the County's Economic Engine

Presented by Robel Worku  
Economic Development Specialist

Montgomery County Economic  
Development Corporation

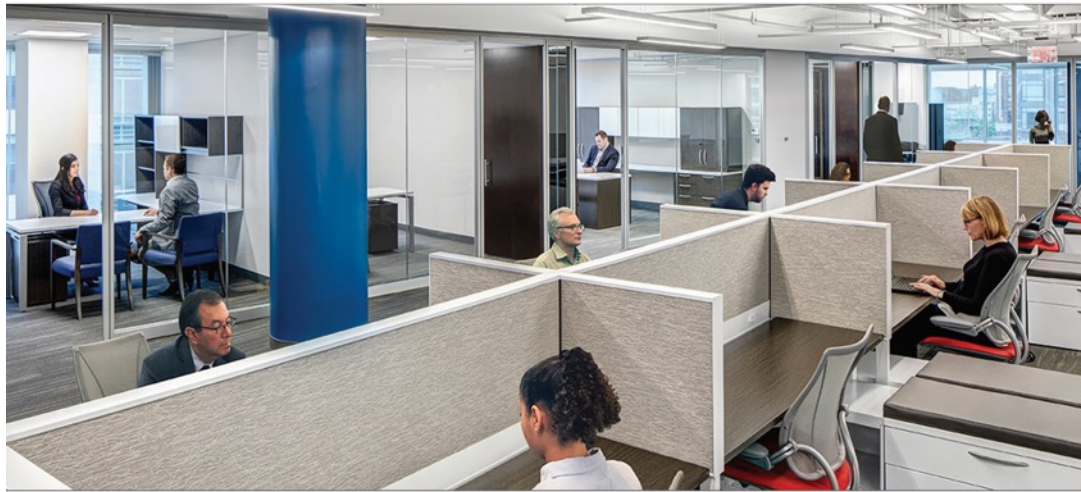
May 18th, 2022



# About MCEDC

The **official public-private economic development organization** representing Montgomery County, MD

Led by a board of directors, our mission is to **help businesses start and grow in the county, or help companies relocate here**



**How does MCEDC help?**

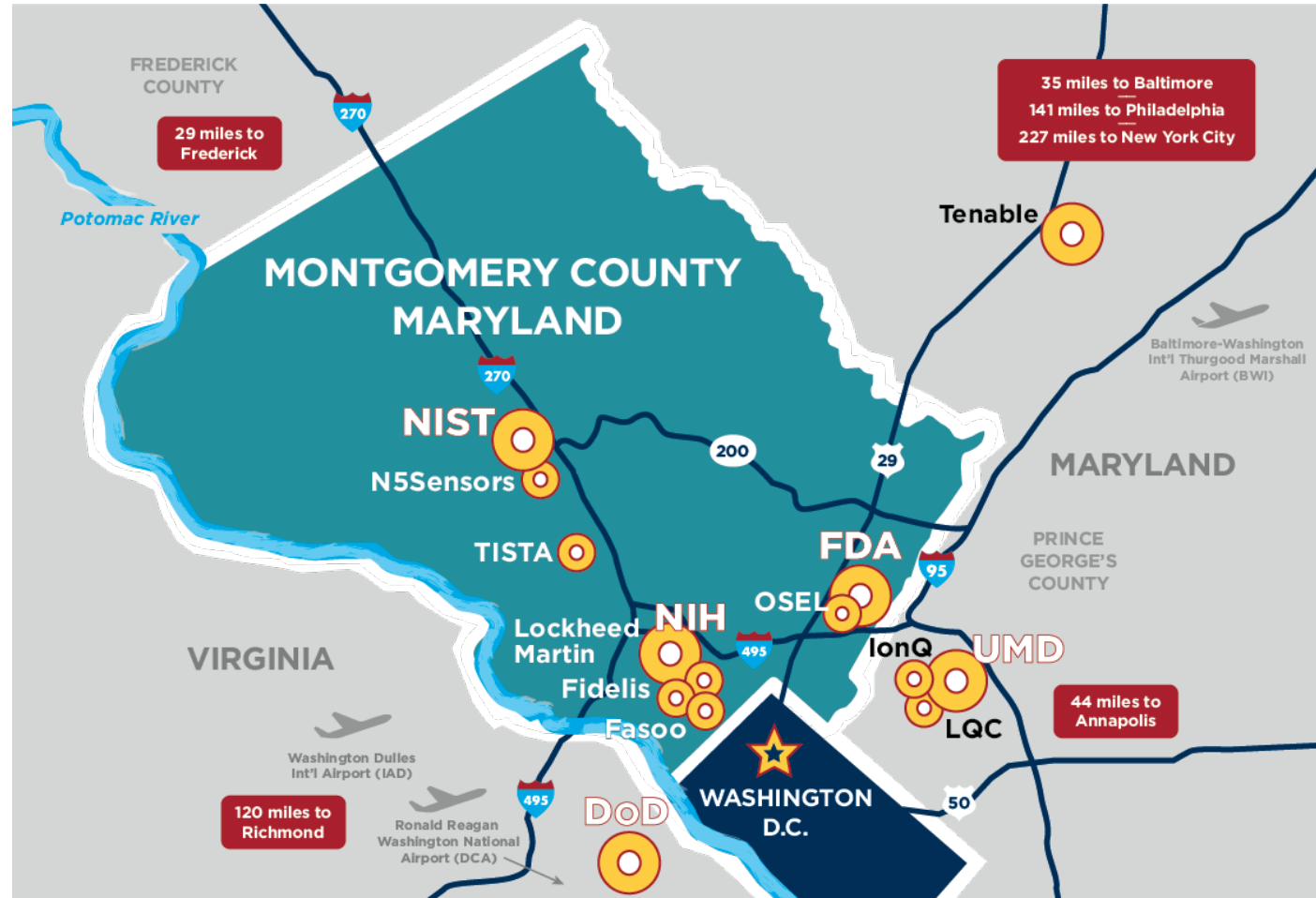
**We help make connections to:**

- Gain market intelligence
- Link business owners to aligned partnerships
- Find the ideal business address
- Explore available incentives
- Attract talent and help with workforce training
- Help companies relocate here

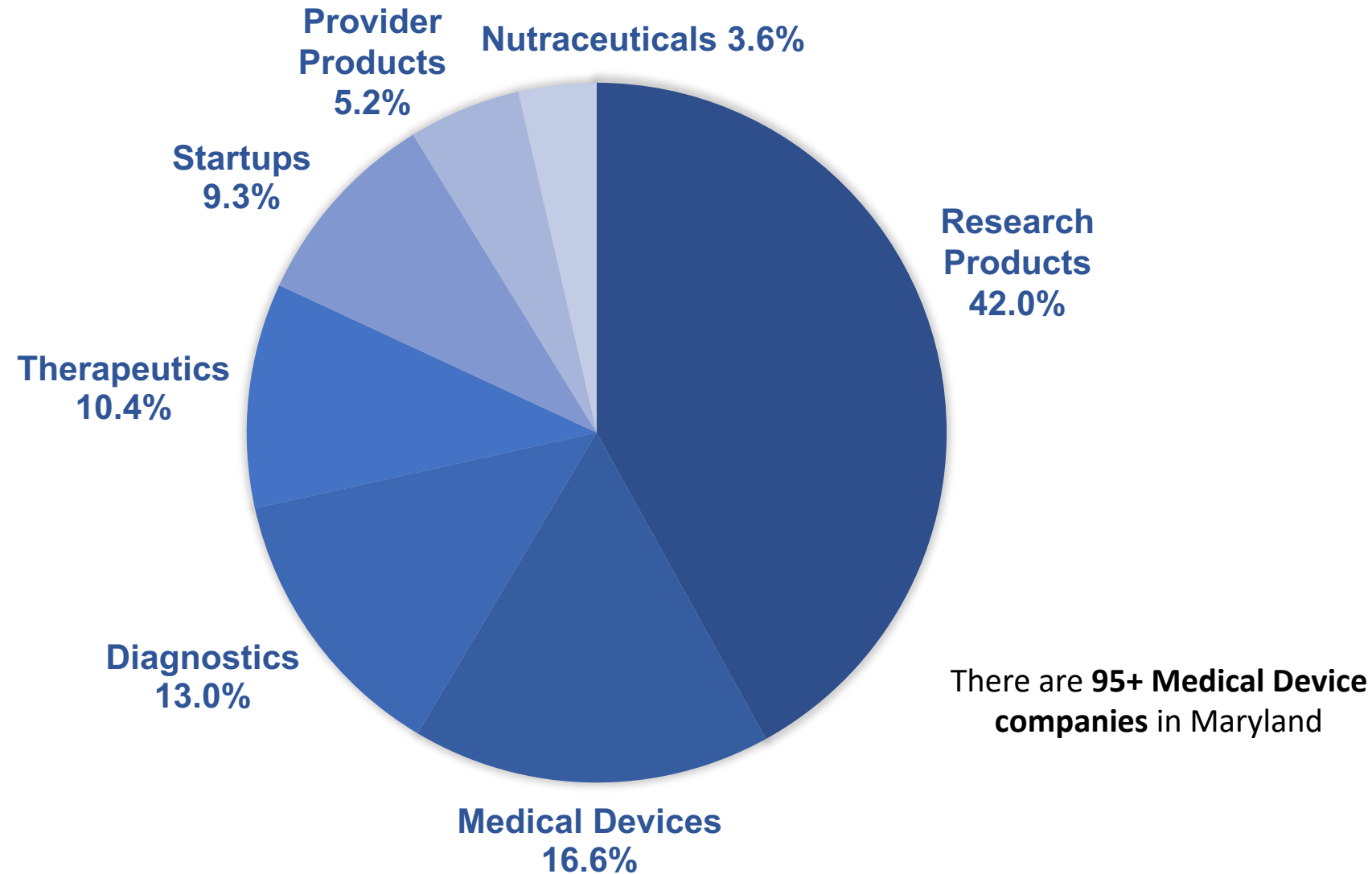


# Big Data Capital Next to the Nation's Capital

Partial list of federal assets and local companies



# Distribution of 800+ BioHealth Companies in Montgomery County





# MONTGOMERY COUNTY LIFE SCIENCES INVESTMENTS

Over **\$1 billion** in Venture Capital raised since 2015

Over **\$245 billion** market cap of companies with  
global or U.S. headquarters in Montgomery County



# Examples of companies using Big Data

## COMPANY USING SUPERCOMPUTERS IN DRUG DISCOVERY

[Gain Therapeutics](#) is a preclinical biotechnology company focused on developing new medicines for protein misfolding diseases. Gain's supercomputer-driven, target-based drug discovery platform, "SEE-Tx™," is a novel approach that uses a convergence of Computational Biology and Supercomputing to discover previously unidentified allosteric binding sites.

## COMPANY LOOKING FOR ADVANCED COMPUTATIONAL METHODS AND HARDWARE TO SOLVE BIG DATA PROBLEMS IN BIO

[EzBiome](#) The Gaithersburg company uses data and databases-driven precision taxonomy with the world's largest curated reference database, and advanced computational platform built-on cutting-edge genomic intelligence.



# Helping in Pandemic Recovery Efforts

During the pandemic, in addition to ongoing business support, MCEDC has pivoted to help the economic recovery for our businesses. Support grant programs include:

**Bio Lab Pilot** – \$96,000 grant program to six small life science companies (administered pre-pandemic)

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**Restaurant Relief Fund** – 3 rounds of relief for a total of \$16.5M dollars distributed to over 900 restaurants

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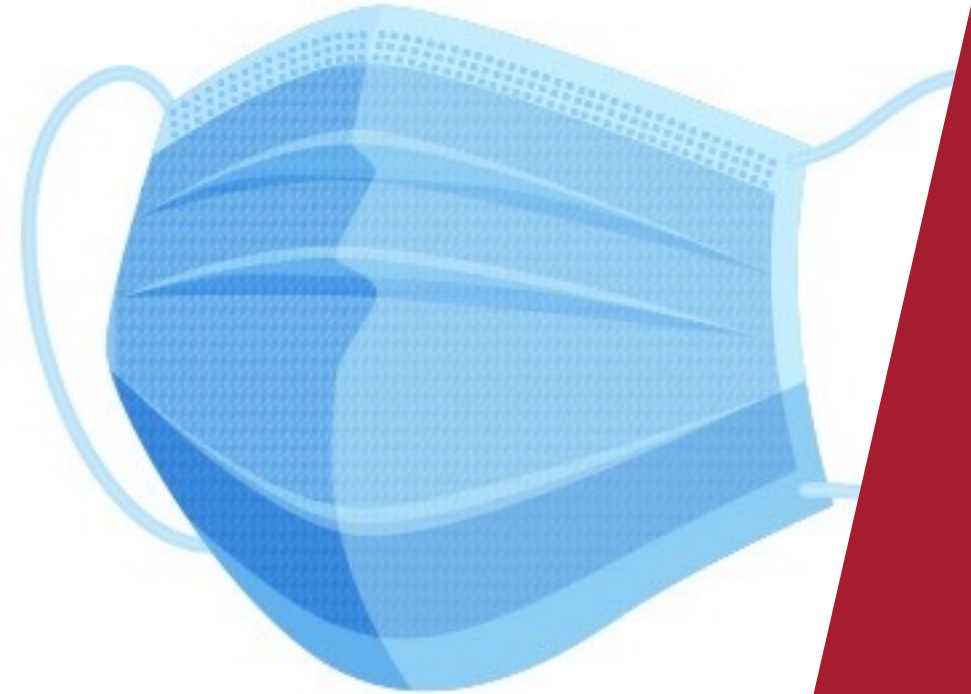
**Telework Fund** – \$1M to help businesses purchase telework equipment and software

---

**Local Production Fund** – \$200,000 fund to help businesses pivot to making Personal Protection Equipment (PPE) to support local pandemic needs

---

**3R Initiative Restaurant & Retail Grants** – \$1M program to assist restaurants and retailers before the 2020 holiday season and winter



# Major Industries

[thinkmoco.com/key-industries](http://thinkmoco.com/key-industries)

BioHealth and Life Sciences

Cybersecurity

Tech & Quantum Computing

Advanced Manufacturing

Hospitality & Tourism

Financial Services

Agribusiness

Nonprofits

Click on images to download

MONTGOMERY COUNTY, MARYLAND (MoCo)

## THE IMMUNIZATION CAPITAL NEXT TO THE NATION'S CAPITAL

**FUNDING MAGNET**

**\$365 Billion** market cap of companies with global or U.S. HQs in MoCo

**\$8 Billion** in 2020 funding secured by MoCo Bio companies

**THE POWER OF OUR FEDERAL PRESENCE**

**\$1 Trillion** annual budget Centers for Medicare and Medicaid Services (CMS) nearby in Baltimore, MD

**\$41 Billion** annual budget National Institutes of Health (NIH)

**\$5 Billion** annual budget Food and Drug Administration (FDA)

**18 Federal Agency Headquarters** located in Montgomery County, Maryland

**\$1.6 Trillion** annual spend from the Hub of Global Healthcare

**\$3 Billion+** invested in MoCo biotech companies for coronavirus vaccine development and production

**THE TALENT CAPITAL**

**300+** Bio companies

**40,000** life science workers

**59%** of adults over 25 have a Bachelor's Degree or higher

**#1** Maryland has the highest concentration of STEM jobs of any state in the U.S.

**#2** Maryland is one of the top states for professional and technical workers

**31.8%** of adults over 25 with a Master's Degree or higher (highest of all counties with more than 1M residents)

Reach out to us to grow in Montgomery County, Maryland  
[connect@thinkmoco.com](mailto:connect@thinkmoco.com)

**MONTGOMERY COUNTY ECONOMIC DEVELOPMENT CORPORATION MARYLAND**

1801 Rockville Pike, Suite 320, Rockville, MD 20852 | 240.646.6700 | [thinkmoco.com](http://thinkmoco.com)

## CYBER INNOVATION THRIVES HERE

MONTGOMERY COUNTY, MARYLAND

Grow with us in the region that's home to the nation's top security agencies and powerful intelligence communities. Draw from our top talent. Maryland is #1 in concentration of STEM jobs and one of the highest density locations for cyber engineers. We have the resources, technology and network to help your company serve federal institutions and private companies.

Our prime location is central to NSA, DIA, DoD and numerous other federal and military institutions. We're a top state for cyber talent. It's all here next to the nation's capital. Montgomery County, Maryland.

**#1 STEM job concentration**

**#1 share of all high-tech businesses**

**17** certified NSA/DHS CASE

**\$558M** VC cyber investment in '20

Top names in cybersecurity call Maryland their home in 2020

**ACCESS OUR FEDERAL ASSETS**

**IN MONTGOMERY COUNTY**

- 18 Federal Agency HQs including NSF, NIH & FDA
- 38 Federal Labs
- The National Cybersecurity Center of Excellence (NCCOE)

**IN GREATER MD, DC AND VA**

- 60 Federal Government Agencies
- The U.S. Cyber Command Center
- The National Security Agency
- The Central Intelligence Agency
- 34 Federal Labs
- 20 Military Facilities

**FEDERAL SPENDING**

- \$7.4 Billion+ Cyber-related activities
- \$8.6 Billion+ Support in Dept. of Defense
- Gain priority to major federal agencies

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MONTGOMERY COUNTY, MARYLAND

## THE CONVERGENCE OF TECH, DATA & TALENT

Accelerate Data Center Innovation in Montgomery County, Maryland

Become a part of an exciting region that synchronizes tech, data and talent. Montgomery County is the perfect landing spot for Data Center growth and expansion. Grow here to centralize shared IT operations and equipment for data storage. Montgomery County is ideally prepared to house these critical assets vital to smart cities and daily operations for business.

**Why Montgomery County? Find key assets for growth:**

- Rich in the talent you need to successfully run the data center
- Ideal and cool location which houses the National Cybersecurity Center of Excellence (NCCOE), a part of the National Institute of Standards and Technology (NIST), along with 18 federal agency headquarters
- Private company tech excellence: 1,200 tech firms and more than 80,000 tech workers in the county
- ultraMontgomery supports cost-effective, competitive access to robust, reliable and secure broadband services and ultra-high speed reliability for businesses throughout Montgomery County

**MONTGOMERY COUNTY VS. THE NATIONAL AVERAGE**

Selected Occupation	Employment Numbers	Unemployment Quotient
Software Engineers	801	6.42
Computer and Mathematical Occupations	29,868	1.84
Computer and Information Research Scientists	719	3.91
Information Security Analysts	1,140	2.79
Computer Network Architects	1,207	2.87
Network and Computer Systems Administrators	2,759	2.35
Mathematicians	47	4.67
Operations Research Analysts	834	3.87

Source: BLSdata 12/2020

**#1** MD has the highest concentration of STEM jobs in the U.S.

**3 times** more cyber-related programs in Maryland than the rest of the country, combined

**29,868** Computer and Mathematical professionals in Montgomery County

**3 1/2 times** more cyber engineers in MD/DC/VA than the rest of the country, combined

Montgomery County and the State of Maryland have the infrastructure and innovation to grow your Data Center. [thinkmoco.com](http://thinkmoco.com)

MONTGOMERY COUNTY, MARYLAND

## THE VACCINE CAPITAL NEXT TO THE NATION'S CAPITAL

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THANK YOU

Visit us at [thinkmoco.com](http://thinkmoco.com)

[robelt@thinkmoco.com](mailto:robelt@thinkmoco.com)

Sign up for our [newsletter](#) for ongoing business news and support  
Send us your updates so we can help promote your business — email us at [connect@thinkmoco.com](mailto:connect@thinkmoco.com)

# Workshop Overview

Ron Pulivarti, NIST NCCoE

# WHO WE ARE



A **solution-driven, collaborative** hub addressing complex cybersecurity problems



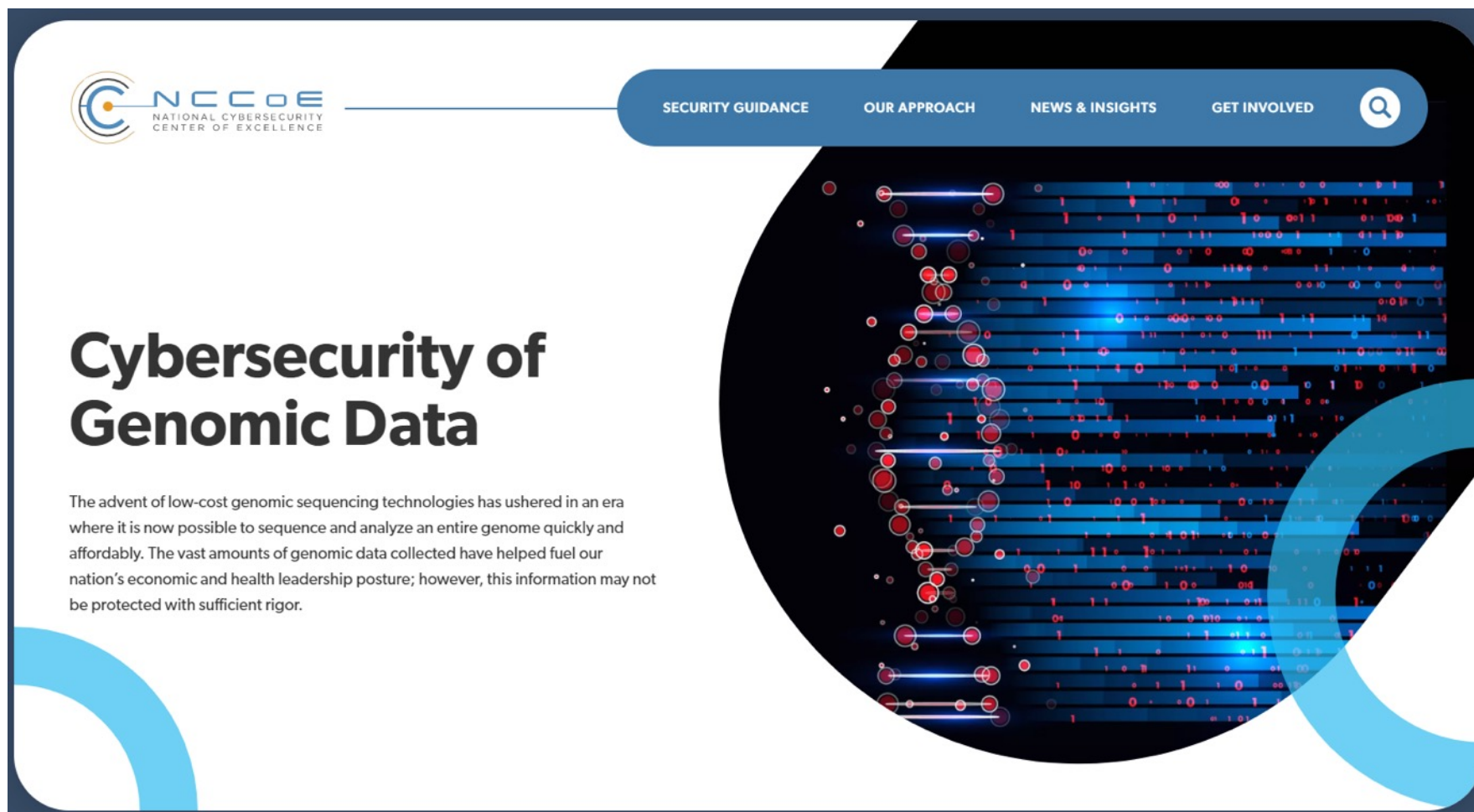


# DISCLAIMER



Certain commercial entities, equipment, products, or materials may be identified by name or company logo or other insignia in order to acknowledge their participation in this collaboration or to describe a procedure or concept adequately. Such identification is not intended to imply special status or relationship with NIST or recommendation or endorsement by NIST or NCCoE.

# PAST WORK: NCCOE VIRTUAL WORKSHOP ON THE CYBERSECURITY OF GENOMIC DATA



<https://www.nccoe.nist.gov/projects/cybersecurity-genomic-data>

# AGENDA: MAY 18



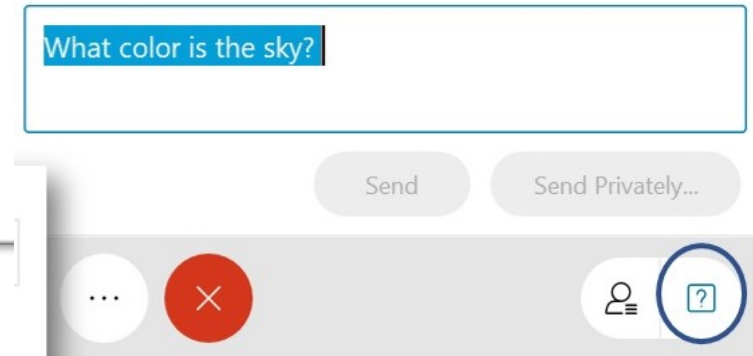
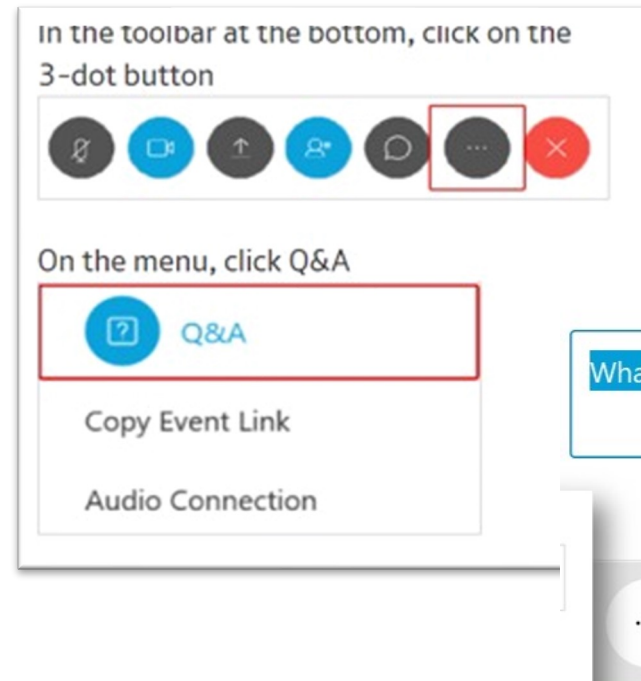
<i>Segment</i>	<i>Time (EDT)</i>
Workshop Overview	1:00 PM – 1:15 PM
Session One: Genomic Sequencer Device Security	1:15 PM – 2:15 PM
Break	2:15 PM – 2:30 PM
Session Two: Securing Sensitive Human Data in support of Genomics Research	2:30 PM – 3:25 PM
Wrap Up	3:25 PM – 3:30 PM



# Audience Engagement

Please use the Q&A window to enter your questions for today's workshop. We will do our best to answer the questions in real time.

1. On the right side, click on Q&A header to open the Q&A panel.
2. Type your question in the box, along with your name and organization.
3. Click **send**.
4. We will answer as many questions as we are able during Q&A sessions.



# Housekeeping

- We support the health and well being for all.
  - We are supporting virtual collaboration.
  - We have a 15-minute break planned for the day.
- We want audience engagement.
  - Please pose your questions for today's workshop using the Q&A window.
- We intend to share our learnings today.
  - We are recording this session for future post on the NCCoE Website.
  - We will post the speaker slides and recording on the NCCoE Website.

**This meeting is  
being recorded.**

# Genomic Sequencer Device Security

Paul Watrobski (NIST)

Blaine Mulugeta (MITRE)

Charles Fracchia (BIO-ISAC)

Phillip Whitlow (HudsonAlpha)



# Session Overview

<i>Segment</i>	<i>Time</i>
Overview and Introduction of Panelists	20 minutes
NIST/NCCoE IoT/MUD Project Overview	
Bio-ISAC Introduction	
HudsonAlpha Introduction	
Panel Discussion <ul style="list-style-type: none"><li>• Current approaches to bio-security of devices</li><li>• Future use of MUD for sequencers</li><li>• Future need for sequencer security baselines</li></ul>	30 minutes
Q&A	10 minutes

# Cybersecurity Challenges

## Securing Genetic Sequencers

What's driving the need for solutions?

Sequencers are essentially IoT devices connected to the enterprise network

- Accessible and remotely managed
- Connected systems and infrastructure
- Proprietary hardware and software
- Limited configuration standards
- Increased complexity and data volumes



# Panelists

Paul Watrobski – NIST

Blaine Mulugeta – MITRE

Charles Fracchia – BIO-ISAC

Phillip Whitlow – HudsonAlpha



# National Cybersecurity Center of Excellence

## NCCoE Virtual Workshop on Exploring Solutions for the Cybersecurity of Genomic Data

- Paul Watrobski (NIST)
- Blaine Mulugeta (MITRE)

# What is/Why MUD?

# MITIGATING NETWORK-BASED ATTACK USING MUD



## Challenge

- Internet connected devices that are not intended to be used for general purpose computing tasks and have a very specific purpose aren't always capable of protecting themselves (e.g., Genomic Sequencers, IoT devices, etc.)
- Device security may not be a priority due to processing, timing, memory, and power constraints
- Typically, these devices are given full connectivity to the internet by default
- Networked devices can be detected within minutes and exploited due to known security flaws, leading to easily scalable attacks

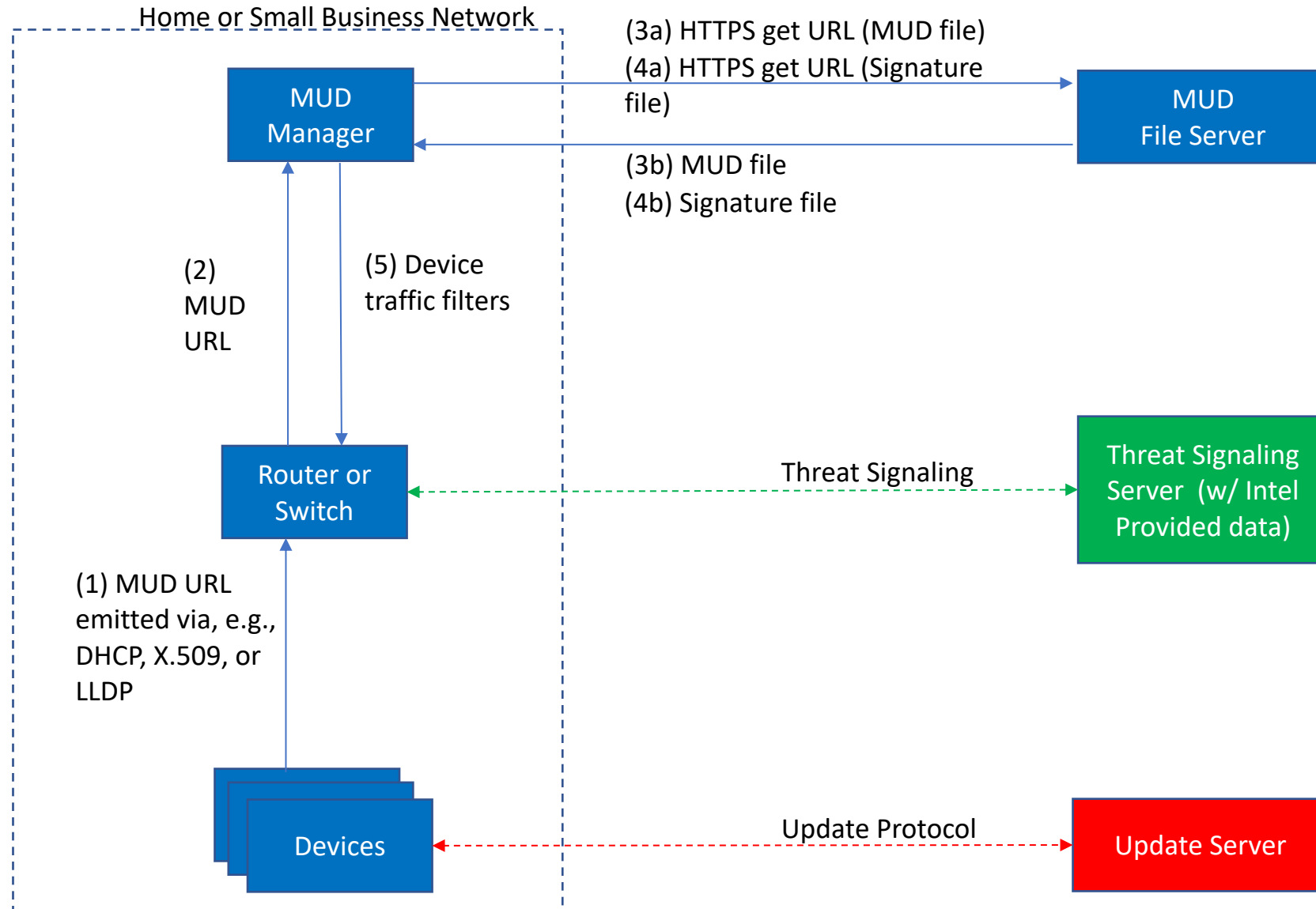
## Solution

- Since these devices have a small number of communication patterns which follows from those small number of uses, the combination of these two statements has been coined Manufacturer Usage Description (MUD) that can be applied at various points within a network.
- MUD enables a network to limit a device's communication within the local network and externally to only those needed and intended by the manufacturer's design.
- MUD primarily addresses threats to the device rather than treating the device as a threat
- Use network gateway components and security-aware devices that leverage the MUD Specification (RFC 8520)

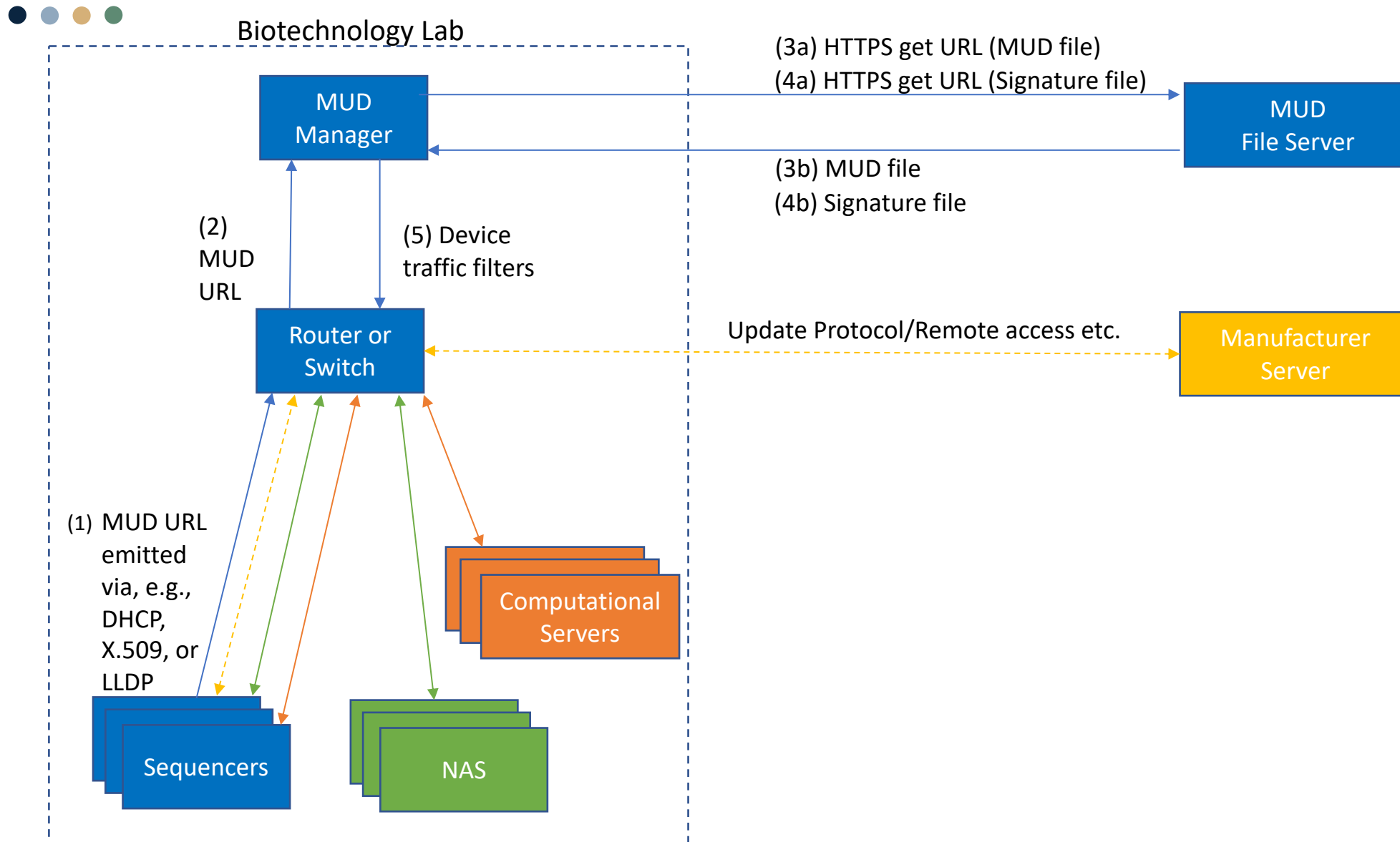
# Architecture Overview



# MITIGATING NETWORK-BASED ATTACKS USING MUD - REFERENCE ARCHITECTURE



# SEQUENCER EXAMPLE ARCHITECTURE





# BIO-ISAC

## Securing the bioeconomy

Charles Fracchia

NIST NCCOE  
18 MAY 2022





# Overview: BIO-ISAC

**BIO-ISAC** was founded to address the demand & meet the rising needs of the bioeconomy community.

**BIO-ISAC** is an international organization that addresses threats unique to the bioeconomy and enables coordination among stakeholders to facilitate a robust and secure industry. <https://isac.bio>





# **BIO-ISAC Mission - in a nutshell**

**Provide concrete solutions to “all hazards” affecting the bioeconomy.**



# BIO-ISAC Benefit - Free Emergency Threat Hunting



WIRED

SIGN IN



LILY HAY NEWMAN

SECURITY MAY 12, 2022 8:30 AM

## The Hidden Race to Protect the US Bioeconomy From Hacker Threats

A biotech threat intelligence group is gaining supporters as urgency mounts around an overlooked vulnerable sector.



Thank you to founding member

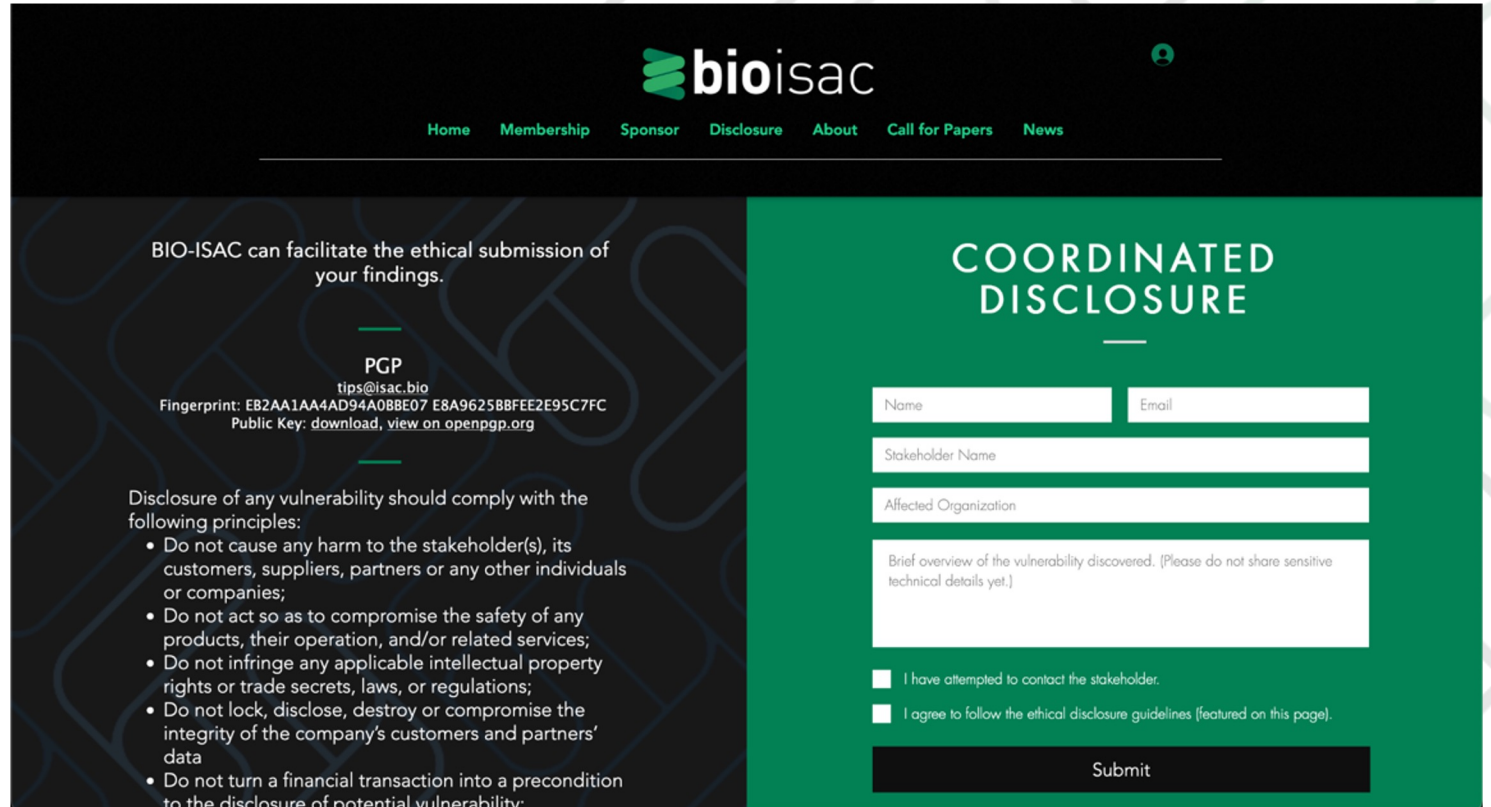


for their support & partnership

# Contribute disclosures

**Have discovered a vulnerability or issue with your bioindustrial, lab, agro, analytical, pre-clinical or other systems?**

BIO-ISAC can help you responsibly disclose the vulnerability and get it addressed.



The screenshot shows the BIO-ISAC website's 'COORDINATED DISCLOSURE' page. The header features the BIO-ISAC logo and navigation links: Home, Membership, Sponsor, Disclosure, About, Call for Papers, and News. The main content area is split into two columns. The left column, on a dark background, explains that BIO-ISAC facilitates ethical submissions and provides PGP contact information for tips@isac.bio, including a fingerprint and public key. It also lists principles for responsible disclosure, such as not causing harm and not compromising safety. The right column, on a green background, contains a form with fields for Name, Email, Stakeholder Name, and Affected Organization. It also includes a text area for a brief overview of the vulnerability and two checkboxes for user consent. A 'Submit' button is at the bottom right.

**BIO-ISAC**

Home Membership Sponsor Disclosure About Call for Papers News

BIO-ISAC can facilitate the ethical submission of your findings.

PGP  
tips@isac.bio  
Fingerprint: EB2AA1AA4AD94A0BBE07 E8A96258BFEE2E95C7FC  
Public Key: download, view on openpgp.org

Disclosure of any vulnerability should comply with the following principles:

- Do not cause any harm to the stakeholder(s), its customers, suppliers, partners or any other individuals or companies;
- Do not act so as to compromise the safety of any products, their operation, and/or related services;
- Do not infringe any applicable intellectual property rights or trade secrets, laws, or regulations;
- Do not lock, disclose, destroy or compromise the integrity of the company's customers and partners' data
- Do not turn a financial transaction into a precondition to the disclosure of potential vulnerability;

**COORDINATED DISCLOSURE**

Name  Email

Stakeholder Name

Affected Organization

Brief overview of the vulnerability discovered. (Please do not share sensitive technical details yet.)

☐ I have attempted to contact the stakeholder.

☐ I agree to follow the ethical disclosure guidelines (featured on this page).

Submit

<https://isac.bio/disclosure>



# Thank you

Prepared for NIST NCCOE  
18 MAY 2022







Phillip Whitlow

# Who is HudsonAlpha?

HudsonAlpha Institute for Biotechnology is a nonprofit institute founded in 2008 and is a national and international leader in genetics and genomics research and biotech education.

Tens of thousands of genomes (human and non-human) sequenced per year on campus

Sequencing use cases include:

- Genetic testing and whole genome sequencing to diagnose patients with rare and/or hard to diagnose genetic illnesses
- Clinical genome sequencing for labs, healthcare institutions, and physicians for diagnostics and treatment of patients
- Developing customized genomic screening programs for patient and employee populations of health systems, physician networks, and self-insured employers
- Original sequencing of plants to facilitate improvements in food production and biofuel research among others

We also host more than 50 associate companies on our campus - all of them involved in bioscience and many performing genomic sequencing in their own labs



# Panel Discussion

Paul Watrobski - NIST

Blaine Mulugeta - MITRE

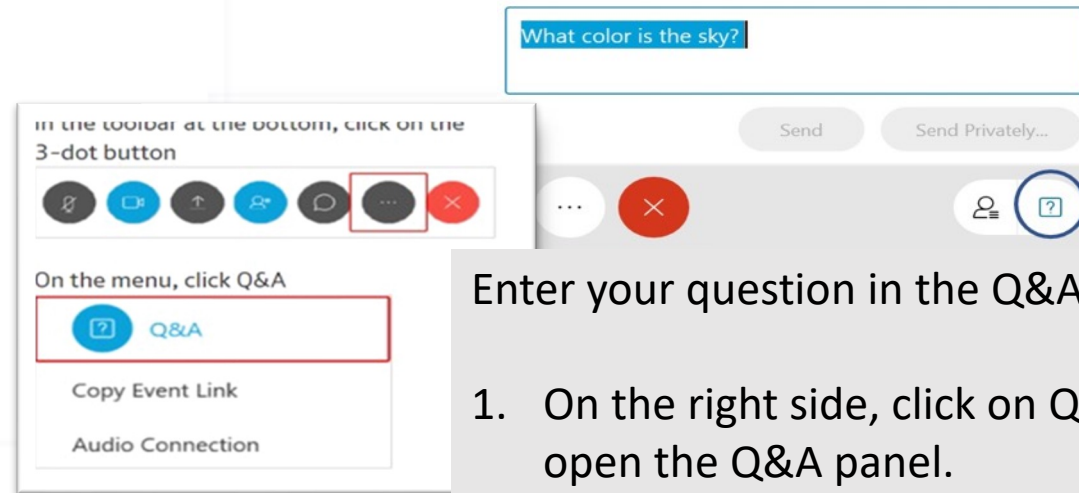
Charles Fracchia - BIO-ISAC

Phillip Whitlow - HudsonAlpha



# Genomic Sequencer Device Security

## Moderated Questions and Answers



Enter your question in the Q&A panel.

1. On the right side, click on Q&A header to open the Q&A panel.
2. Type in the box **your name, organization and question.**
3. Click send.



# Break

Enjoy your break.  
We'll start again soon!

2:30 PM

## Coming up next!

TOPIC	PRESENTERS
Session Two: Securing Sensitive Human Data in support of Genomics Research	Mike Feolo (NIH) Kurt Rodarmer (NIH)

# Welcome Back!

This meeting is being recorded.

# Securing Sensitive Human Data in support of Genomics Research

Mike Feolo (NIH)

Kurt Rodarmer (NIH)

# ***Securing Sensitive Human Data in support of Genomics Research***

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*May 18, 2022*

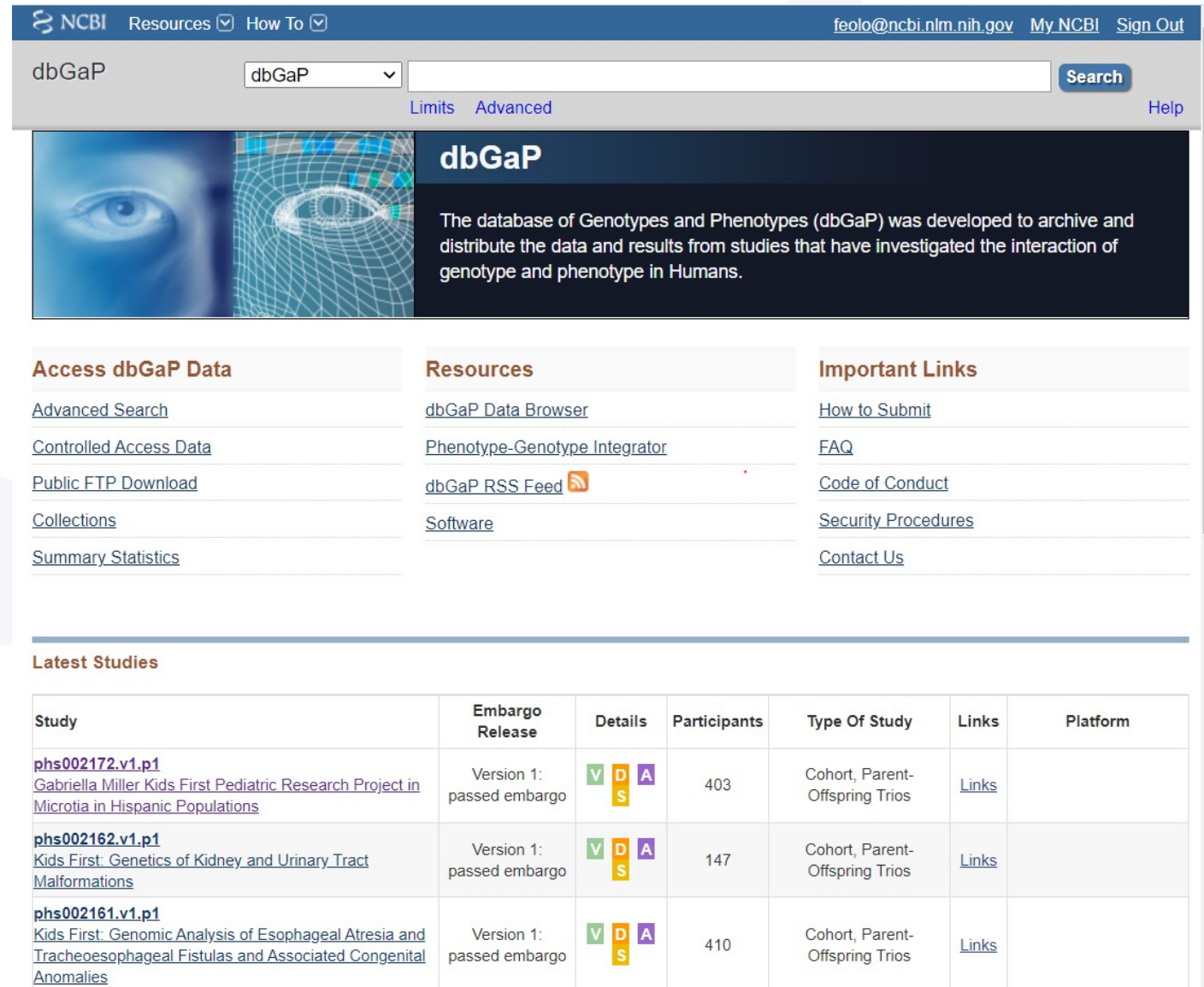
*Michael Feolo  
dbGaP Team Lead  
NIH/NLM/NCBI*

*Kurt W. Rodarmer  
RAS Security Advisory Group  
National Institutes of Health,  
NLM/NCBI*



# dbGaP: The database of Phenotypes and Genotypes

- Permanent Archive
- Versioned Releases
- Public Metadata
- Controlled Access



NCBI Resources How To feolo@ncbi.nlm.nih.gov My NCBI Sign Out

dbGaP dbGaP Search Limits Advanced Help

**dbGaP**

The database of Genotypes and Phenotypes (dbGaP) was developed to archive and distribute the data and results from studies that have investigated the interaction of genotype and phenotype in Humans.

**Access dbGaP Data**

- [Advanced Search](#)
- [Controlled Access Data](#)
- [Public FTP Download](#)
- [Collections](#)
- [Summary Statistics](#)

**Resources**

- [dbGaP Data Browser](#)
- [Phenotype-Genotype Integrator](#)
- [dbGaP RSS Feed](#)
- [Software](#)

**Important Links**

- [How to Submit](#)
- [FAQ](#)
- [Code of Conduct](#)
- [Security Procedures](#)
- [Contact Us](#)

**Latest Studies**

Study	Embargo Release	Details	Participants	Type Of Study	Links	Platform
<a href="#">phs002172.v1.p1</a> <a href="#">Gabriella Miller Kids First Pediatric Research Project in Microtia in Hispanic Populations</a>	Version 1: passed embargo	V D S A	403	Cohort, Parent-Offspring Trios	<a href="#">Links</a>	
<a href="#">phs002162.v1.p1</a> <a href="#">Kids First: Genetics of Kidney and Urinary Tract Malformations</a>	Version 1: passed embargo	V D S A	147	Cohort, Parent-Offspring Trios	<a href="#">Links</a>	
<a href="#">phs002161.v1.p1</a> <a href="#">Kids First: Genomic Analysis of Esophageal Atresia and Tracheoesophageal Fistulas and Associated Congenital Anomalies</a>	Version 1: passed embargo	V D S A	410	Cohort, Parent-Offspring Trios	<a href="#">Links</a>	

# High Level Data Model

## A dbGaP Study

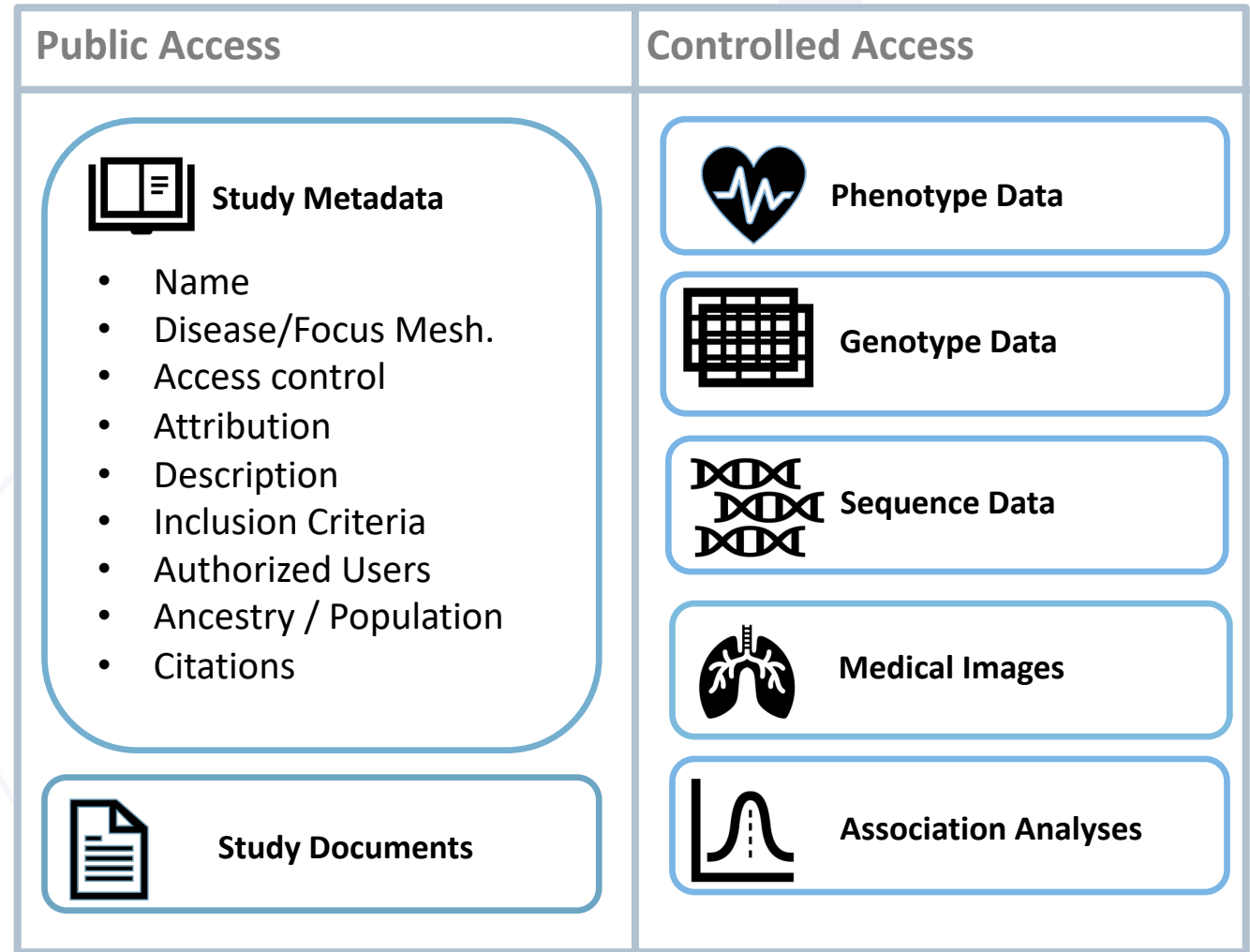
- Data collected from the same funding, Research Focus, Cohort

## Public Access

- Study Level Metadata
- Study Documents
- Summaries and Metadata

## Controlled Access

- Individual Level
- Sensitive Association Analysis



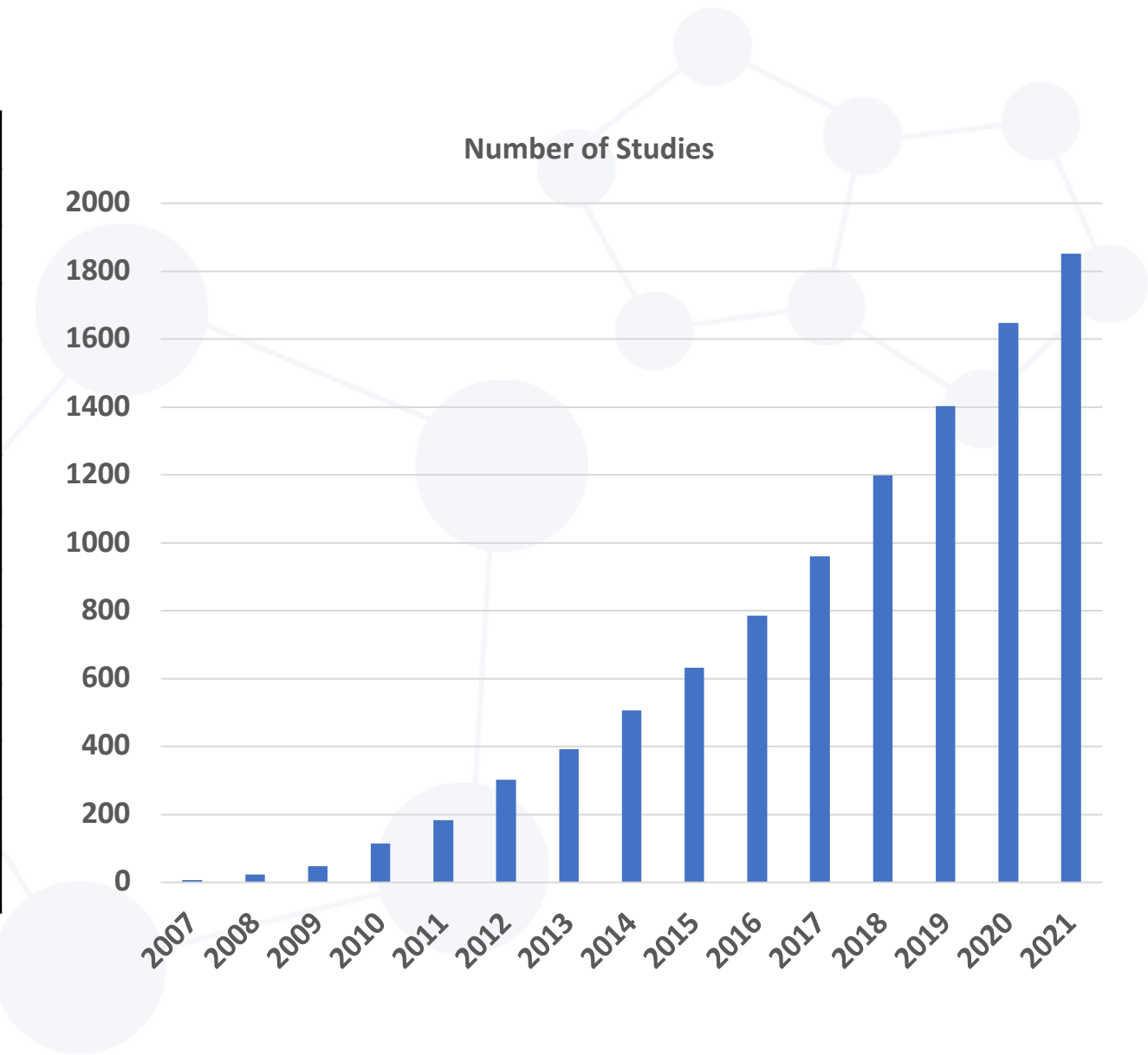
# Why Do this?

- A library of research Data
- Each Study is analogous to a Book
- Scientists from around the world can use these data to augment their research and make new discoveries



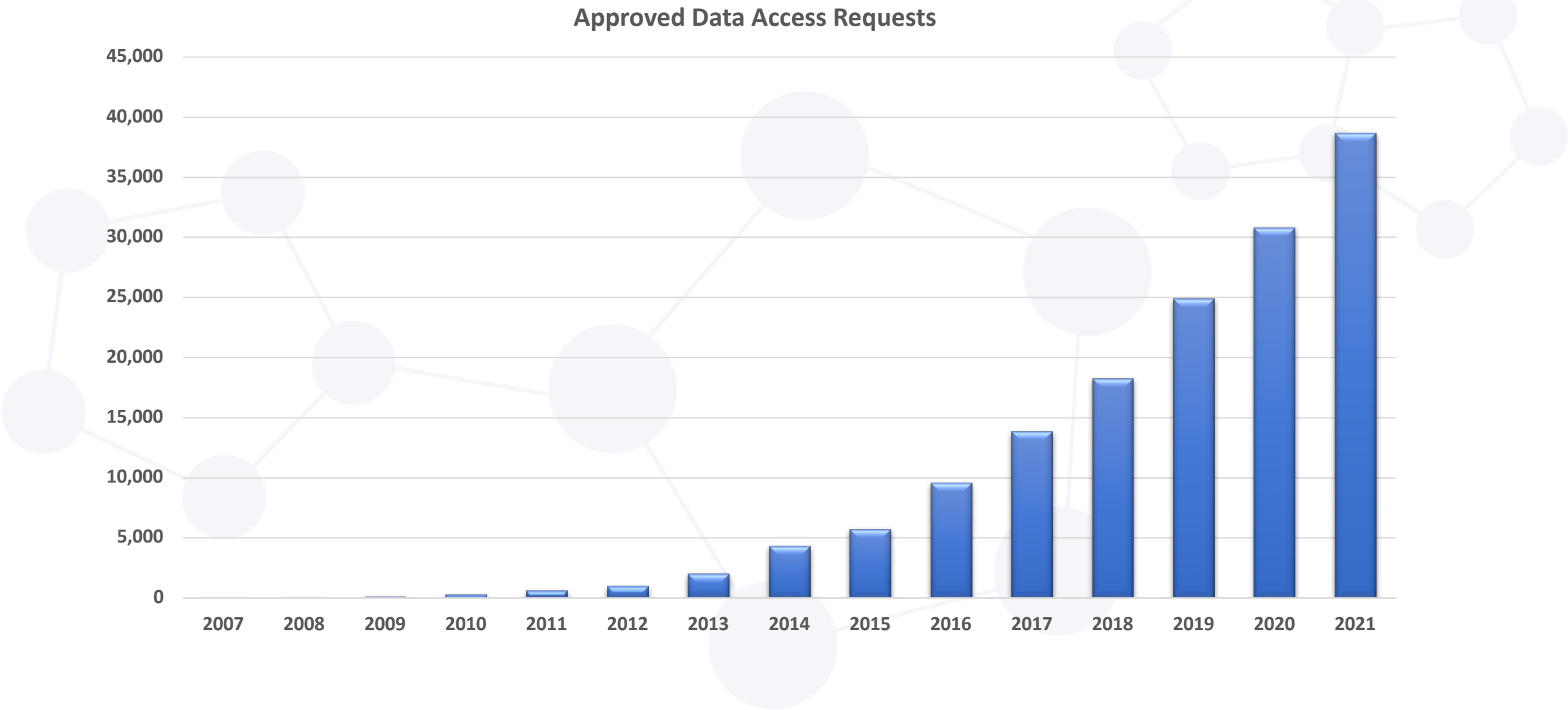
# Data Available in Archive

<b>Studies</b>	1,972
<b>Subjects</b>	~2.9 Million
<b>Samples</b>	~3.4 Million
<b>Phenotype: Variables</b>	371,436
<b>Values</b>	~2.5 Billion
<b>Study Documents</b>	7,149
<b>Association Analyses</b>	7,913
<b>Genotype Assays (array)</b>	~2 Million
<b>Genotype Assays (imputed)</b>	543,137
<b>Genotype Assays (seq derived)</b>	399,269
<b>Sequence (WGS SRA)</b>	178,288
<b>Sequence (WXS SRA)</b>	271,447
<b>Sequence (RNAseq SRA)</b>	86,879
<b>Epigenomic (SRA)</b>	~35,000





# Requests for the data



# Delicate Balance

- Extensive corpus of sensitive human data
- Obligation to shield subject identity
- Value to humanity lies in ability to share within research community
- How to be a resource for the advancement of science while maintaining confidentiality?



# Threats



- Data are stripped of PII
  - But a person's genome is the ultimate PII and is easy to obtain
- Re-identification
  - individuals, families, populations
- Example
  - Politician has a family member within a schizophrenia study

# Threats

## Traditional

- User is traditional threat
  - Legal agreements
  - Institutional liability
  - Penalties for violation





# Threats

## Expanded

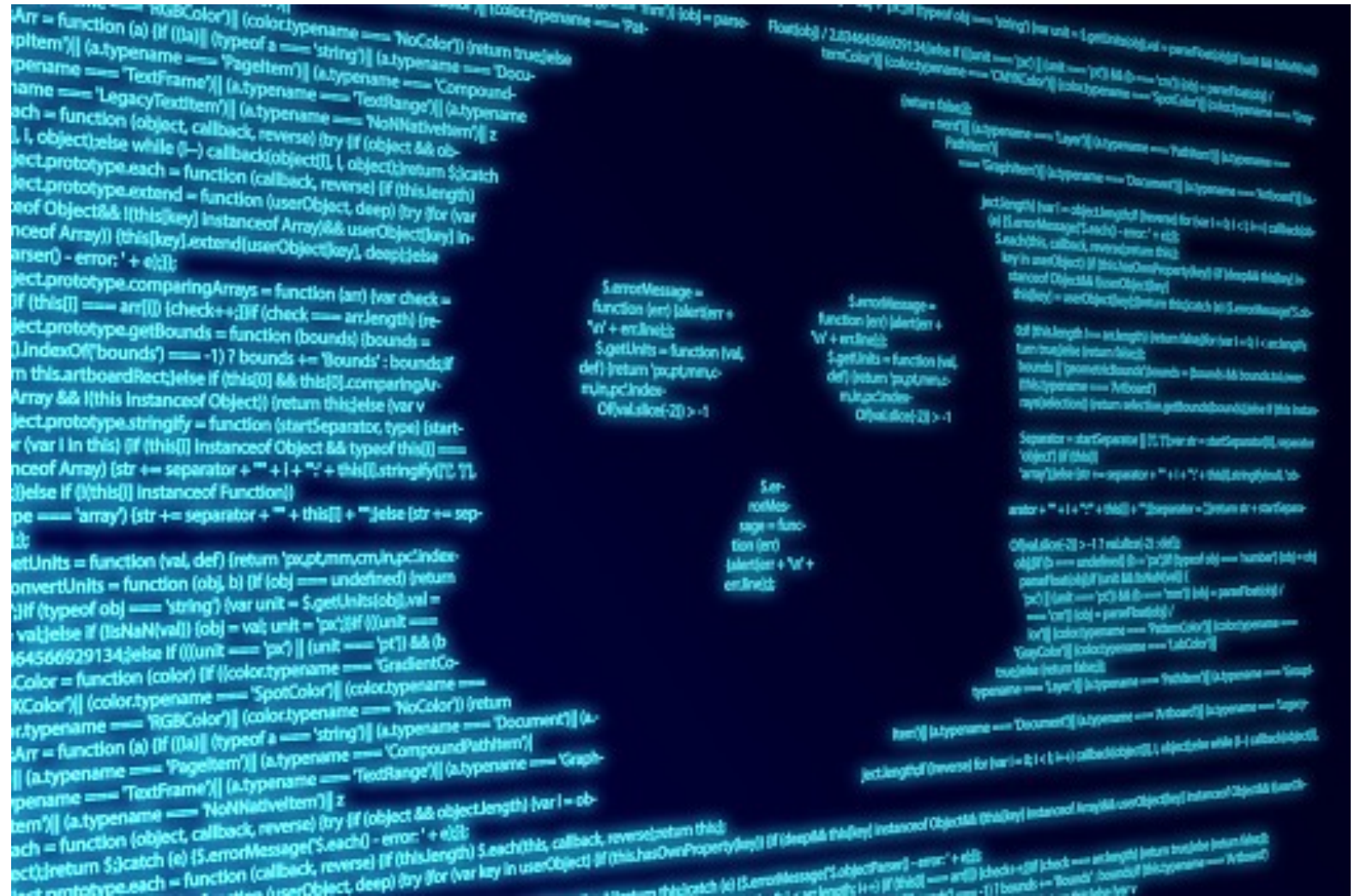


- Intermediate actors present greater threat
  - Systems, networks, clouds
  - Software, workflows
  - None are controlled by user authentication
  - None are affected by legal agreements or penalties

# Threats

## Expanded

- Open-source software chain
  - Tradition of running amateur software
  - All built upon open source of unknown and unverified provenance
  - Few if any pay attention to SBOM



# Threats

## Expanded

- Bringing “compute” to the data
  - Has become a mantra in big data
- Who authorizes the code?
  - Researcher provided
  - Environment provided
- What environments can safely accept unknown code?





# Security Models

- Authority-vs-Identity
  - Traditional identity-based methods prevail
  - NIH is pursuing authority-based models





# Security Models

## Identity-based Security



- Context-free
  - Who you are
- Excess authority
  - All available permissions
- Identity theft
  - Programs use researcher identity
  - Potential for catastrophic consequences
- Poor federation
  - Every system must know every user

# Security Models

## Authority-based Security

- Context-sensitive
  - What you intend to do
- Least authority
  - Only required permissions
- Identity not exploitable
  - Programs restricted to provided permissions
  - Bad or malicious software can be contained
- Good federation
  - Systems recognize source of authority not individual



# Security Models

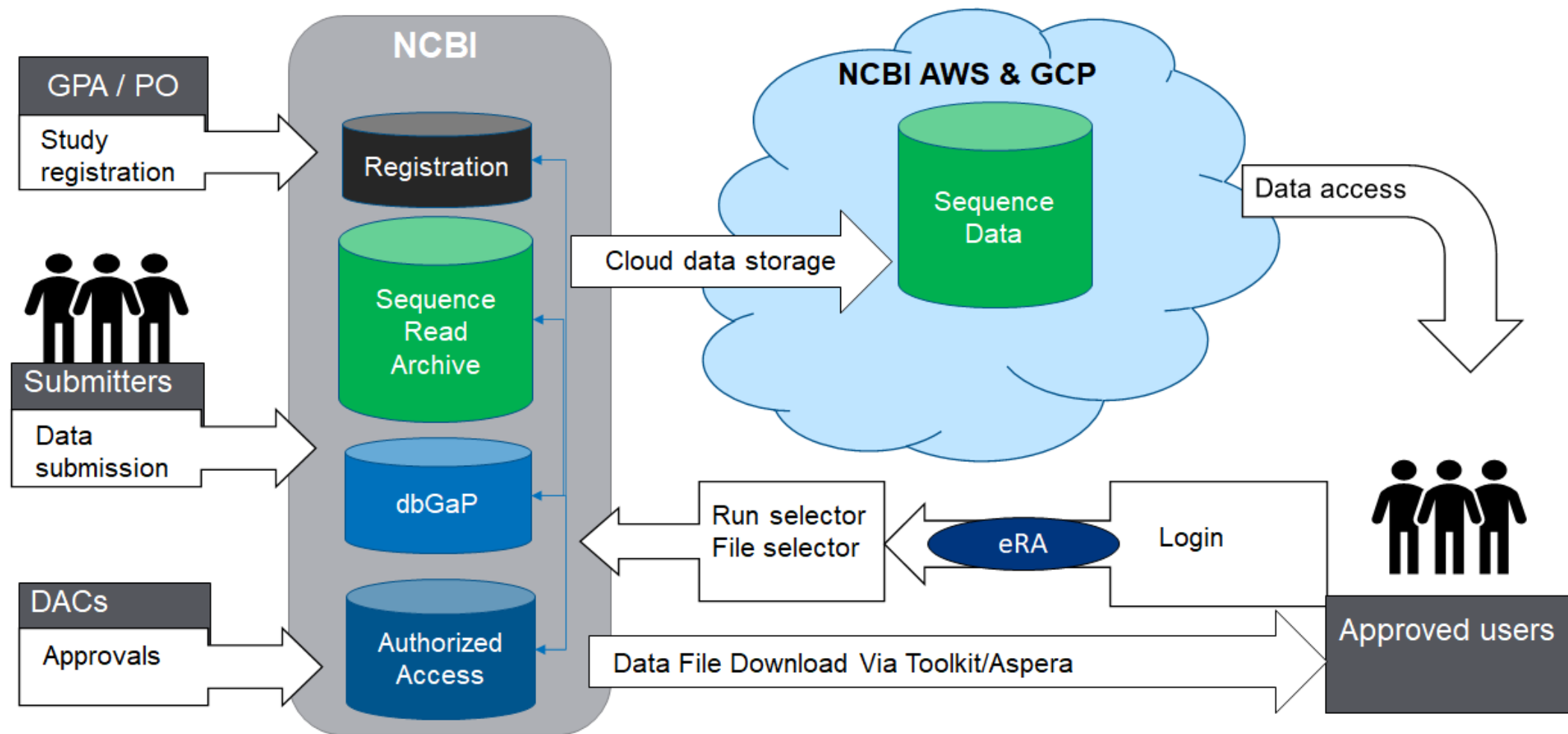
## Compatibility Challenges

- Models do not mix well
- Overwhelming investment in identity-based model



# Journey

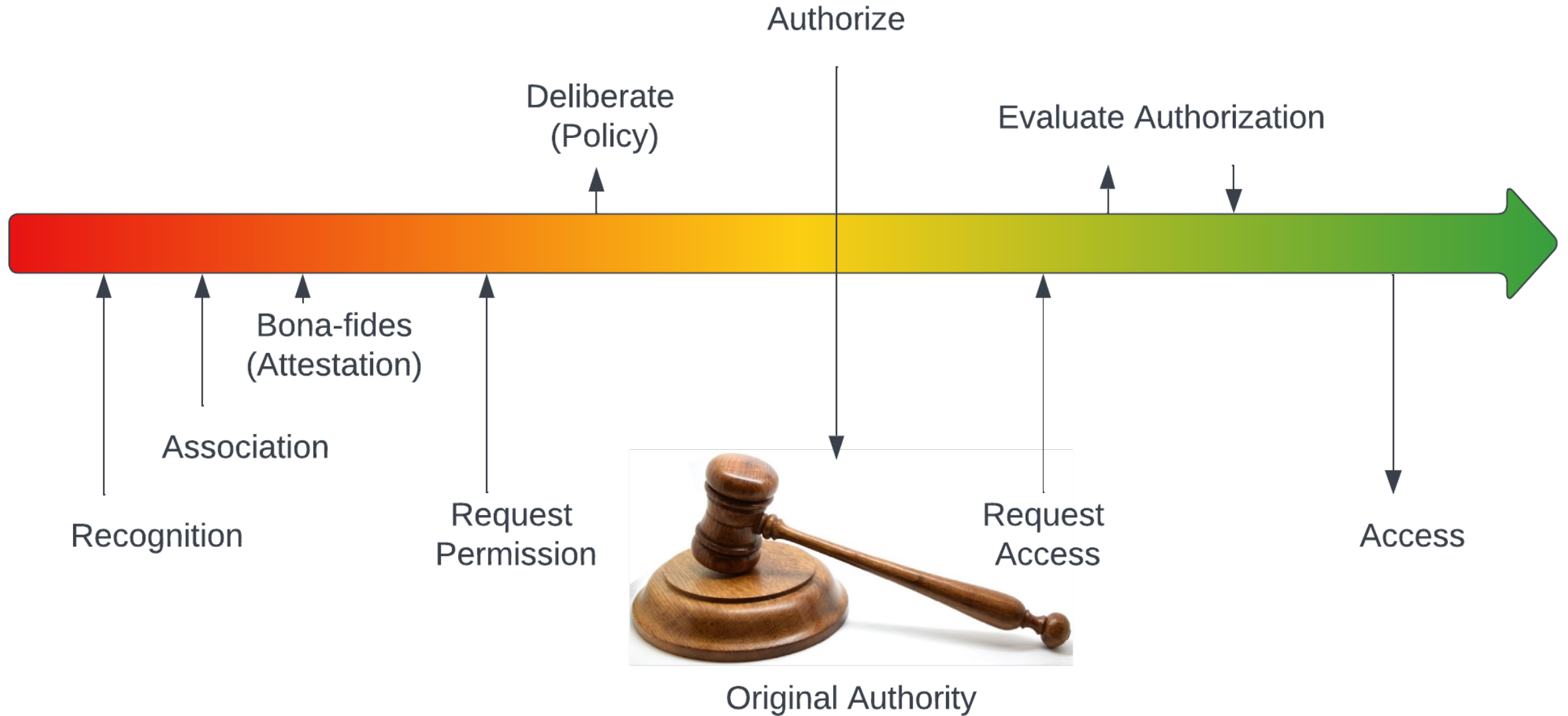
## Submission





# Journey

Access



# Authorization

## Request Access Permissions

- Consent
- Data access request, usage limitations
- Committee approval
- Grant, storage



# Authorization

## Retrieve Permissions

- Grant retrieval
  - Authentication for retrieval of stored authorizations
- NIH-signed Token from RAS
- Token duration
  - Associated with live user



# Computation

## Task Description

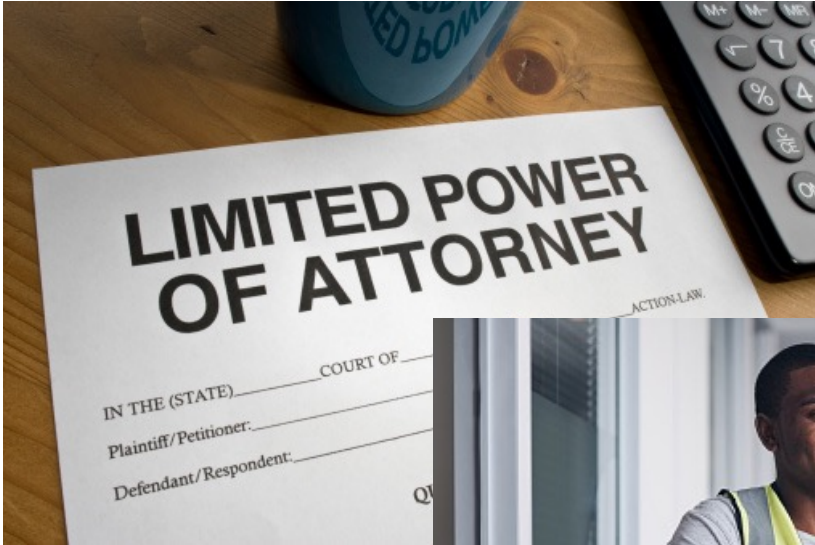
- What to work on
- Least Authority within tokens
- Context-specific limitations





# Computation

## Delegation



- POLA Token
- User not present
- Assign Power of Attorney
  - For program instances
  - For systems
- Worker Authentication
- Token duration



# Zero Trust

Supported by Token Protocol

- System – mTLS
- API – API Token
  - Program to API
  - Bound to system TLS certificate
  - Specific permission subset
- Content – Task-specific Token
  - Delegated user authorizations
  - Assigned to compute environment

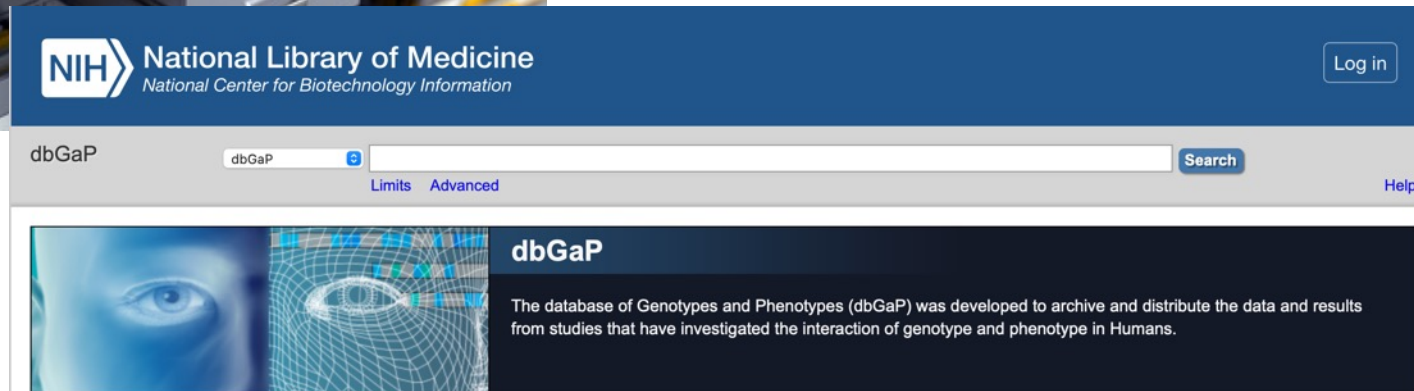


# System Components

## Data Governance Authority



- dbGaP Authorized Access (AA)
  - Data access requests (DAR)
  - Manages authorization process
  - Stores access authorizations
  - Supplies AuthZ to RAS process



# System Components

## Researcher Auth Service (RAS)

- Researcher authentication
  - Anonymized identity
  - 12-hour duration
- Issues signed token
  - Carries authorizations
  - Retrieved from dbGaP
- GA4GH passport
  - Interoperable





# System Components

## Data Locators

- Validates RAS token
- Locate objects by id
- DRS
  - GA4GH emerging standard
  - See: <https://locate.be-md.ncbi.nlm.nih.gov/ga4gh/drs/v1/>
- IDX
  - Convert accessioned object ids to DRS ids
- SDL
  - NCBI SRA established standard





# System Components

## Cloud Storage



- Pre-signed URLs
- Authority vs Identity
  - Require authority-based resource server
  - Employs ACL-based access to objects
- Auth and billing models
  - Unix roots make them inextricably bound
  - Resource servers are billed to provider instead of user
- Surrender of control to CSP
  - Attempts to use CSP IAM bypass tokens

# System Components

## Logging

- Token retrieval at authentication
- Derivation of task-specific token
- Token usage at Data Locator
- Token usage at Resource Server
- Pre-signed URL and bucket logs



# System Components

Audit, Response, Recovery



- Events related by RAS token transaction id
- Raw log ETL and event correlation
- Specific issue detection
- Usage pattern and anomaly detection

# The Confinement Problem

- Prevent leakage after access has occurred
- Accidental leakage
  - Stored insecurely
- Intentional leakage
  - Human relay
- Covert communication
  - Software communications
  - Steganography
- Geographical confinement



# The Confinement Problem

## Genomic Analysis Platform



- Plaintext visibility
- User-provided executables
- Egress
  - Block all egress
  - Filter by content
  - Monitor for violation



# The Confinement Problem

## Secret Store

- No direct visibility
- Predefined executables
  - Difficulties in verification
  - Potential covert channels
- Researcher-defined workflow



# The Confinement Problem

## Cloud Service Providers

- Sponsored projects
  - Created/configured by trusted authority
- Monitored projects
  - Created/configured by user
  - Monitored by trusted authority
- Relies upon cloud IAM
  - May bypass token controls



# Encryption

#011A56AF6420746865206E16700AFFA33C08E00F2A5694C028BE5BF7D011A0010A3BC

32C20736852756B013A00AA206336865206E6741020732C732C20736852756B013A00AA206  
42001AFB71900FF661564207368646F206561A711B2616E642001AFB71900FF661564207368646  
2A5694C028BE5BF7D011A0010A3BCFE561AF87010FC28E00F2A5694C028BE5BF7D011A0010A3BCF  
931001BE45C7710011BFF12FFD9200100D45B333A197716A931001BE45C7710011BFF12FFD920  
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A33C08E00F2A5694C028BE5BF7D011A56AF6101AC9100AFFA33C08E00F2A5694C028BE5BF7D011  
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32C20616E642070617463686513206F5590BF34121000C3732C20616E642070617463686513206  
6206C6974746C6520616E642074616C773192A3BB6C1076C6206C6974746C6520616E642074616  
C2067726173732E2048652061A07072216145A13C7500A16C2067726173732E2048652061A0707  
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420526964696EA120486FAF6420616E013921FC0001016C6420526964696EA120486FAF6420616  
20736865206E61C3AF76656C7920740091AAEF214210656420736865206E61C3AF76656C79207

- Object encryption
- Personalized encryption
- Homomorphic encryption

# Future Directions

- Normalize use of API Tokens
- Greater levels of categorization by sensitivity
- Improve interoperability between CSP and NIH-governed content management





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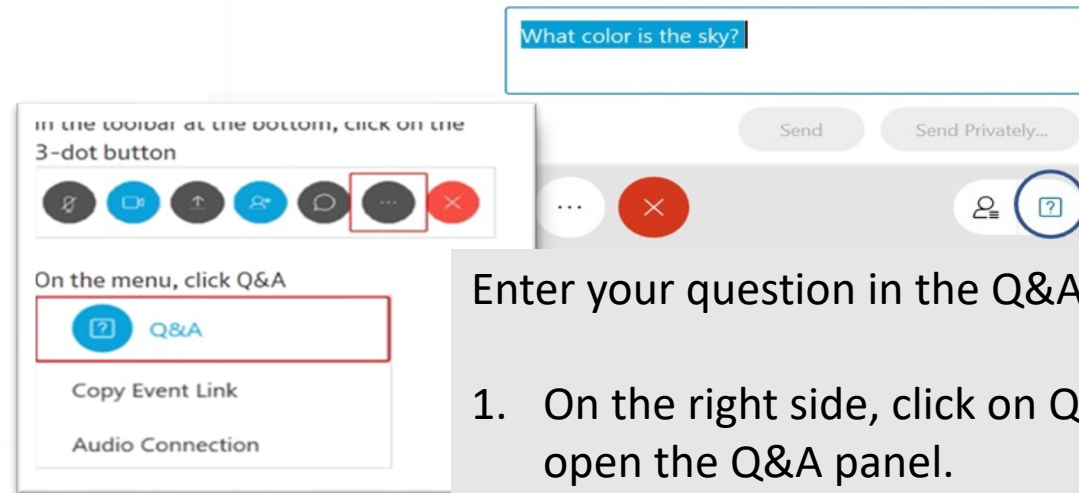
[rodarmer@ncbi.nlm.nih.gov](mailto:rodarmer@ncbi.nlm.nih.gov)

*Thank You!*



# Securing Sensitive Human Data in support of Genomics Research

## Moderated Questions and Answers



Enter your question in the Q&A panel.

1. On the right side, click on Q&A header to open the Q&A panel.
2. Type in the box **your name, organization and question.**
3. Click send.

# Workshop Close Out

Ron Pulivarti, NIST NCCoE



# AGENDA: MAY 19



<i>Segment</i>	<i>Time (EDT)</i>
Workshop Day 1 Reflections	1:00 PM – 1:10 PM
Session Three: Genomic Data Security Through Risk Management	1:10 PM – 2:10 PM
Break	2:10 PM – 2:25 PM
Session Four: Genomic Data Security in Electronic Health Records	2:25 PM – 3:25 PM
Wrap Up	3:25 PM – 3:30 PM

# LINK FOR MAY 19 WORKSHOP



<https://www.nccoe.nist.gov/get-involved/attend-events/nccoe-virtual-workshop-exploring-solutions-cybersecurity-genomic-data>

# Please join us tomorrow!

## Contribute to the conversation

Email [genomic\\_cybersecurity\\_nccoe@nist.gov](mailto:genomic_cybersecurity_nccoe@nist.gov)