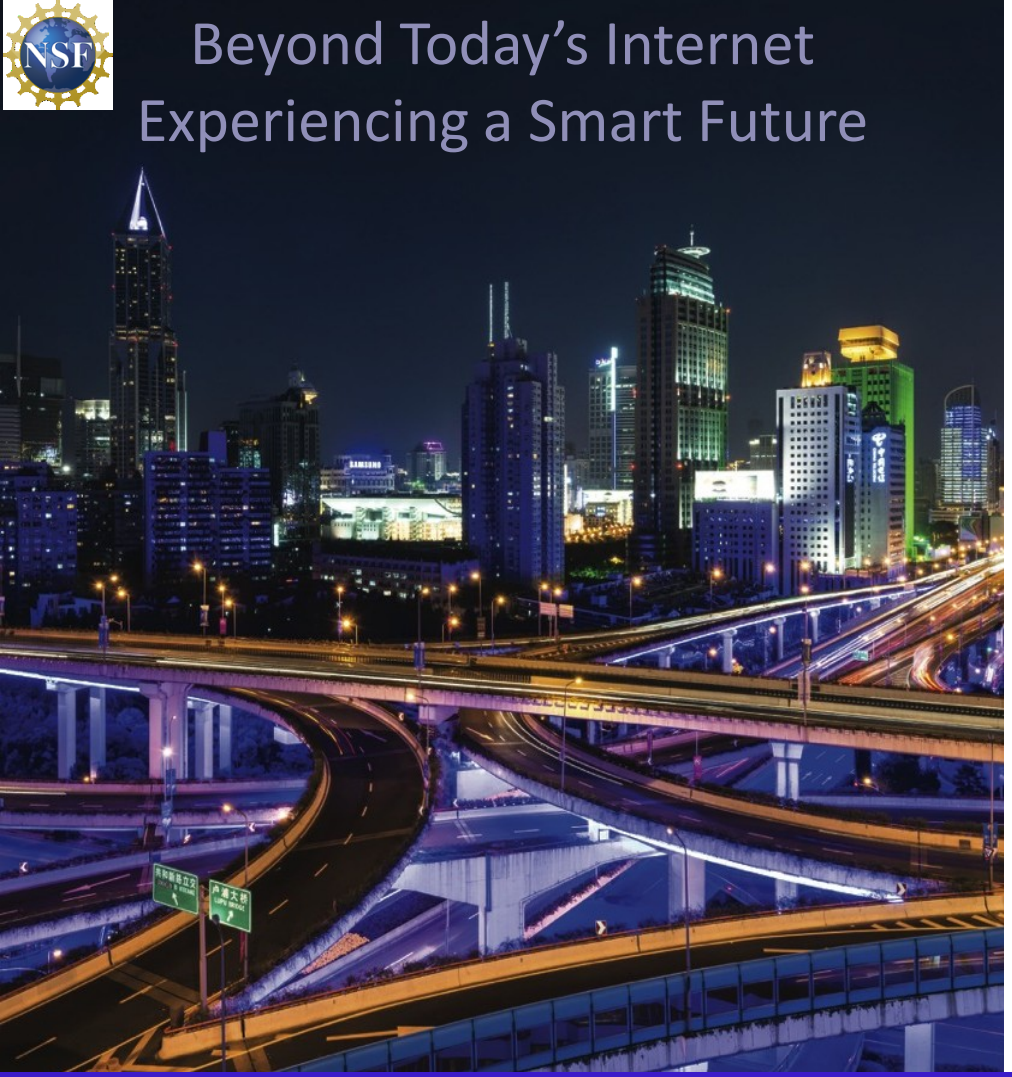


Prototype SDX Bioinformatics Exchange: Demonstrating an Essential Use-Case for Personalized Medicine

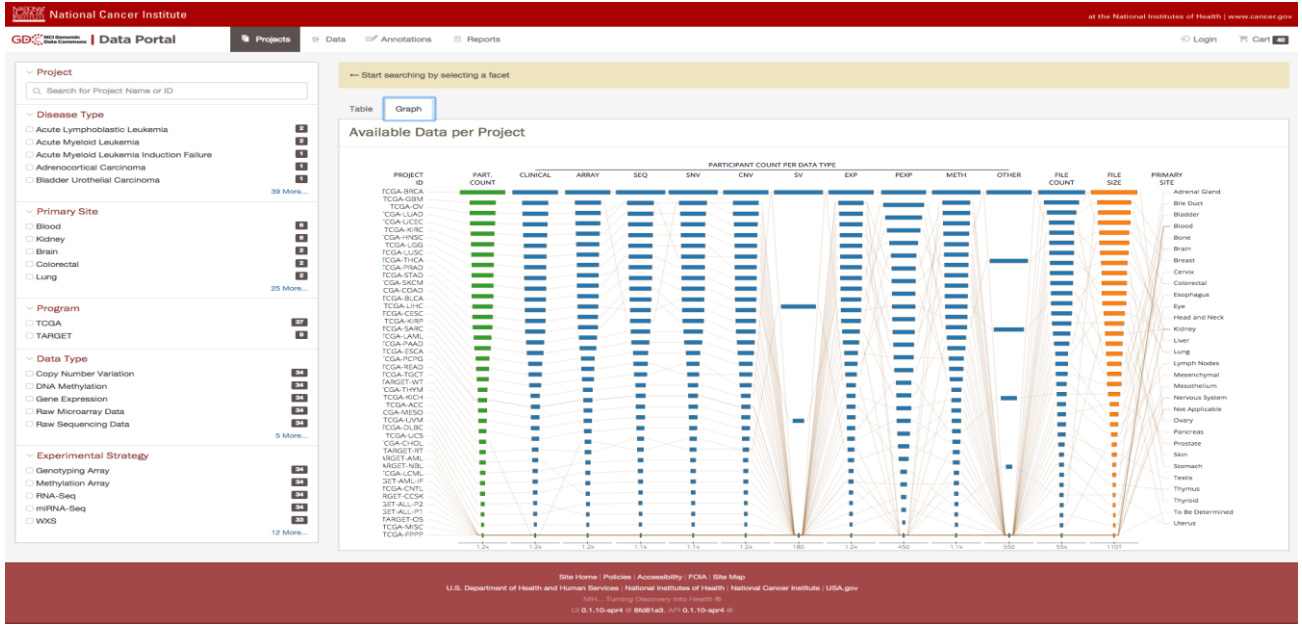
Robert Grossman – University of Chicago
Joe Mambretti – Northwestern University
Piers Nash – University of Chicago
Jim Chen – Northwestern University
Allison Heath – University of Chicago

Precision Medicine

- Precisely match treatments to patients and their specific disease
- Genomic data promises optimal matching.
- 1.7 million cancer cases diagnosed in America each year.
- A single RNA-seq file is 10-20 GB, Whole genome raw data files are > 100 GB.
- Analysis has become the bottleneck and data size is an issue.
  - 2,000,000 genomes ≈ 1 Exabyte (1,000,000,000,000 MB)
  - Cost to sequence 1 genome less than $5,000 and falling fast.
  - Cost to analyze 1 genome is approx. $100,000 and rising.
- A key step towards Algorithm-assisted Personalized medicine is building Data Commons/Cloud analytics and the *Programmable* Networks & Communication Exchanges (SDXs) for high performance, flexible data transport.

Infrastructure for Precision Medicine

NCI Genomic Data Commons

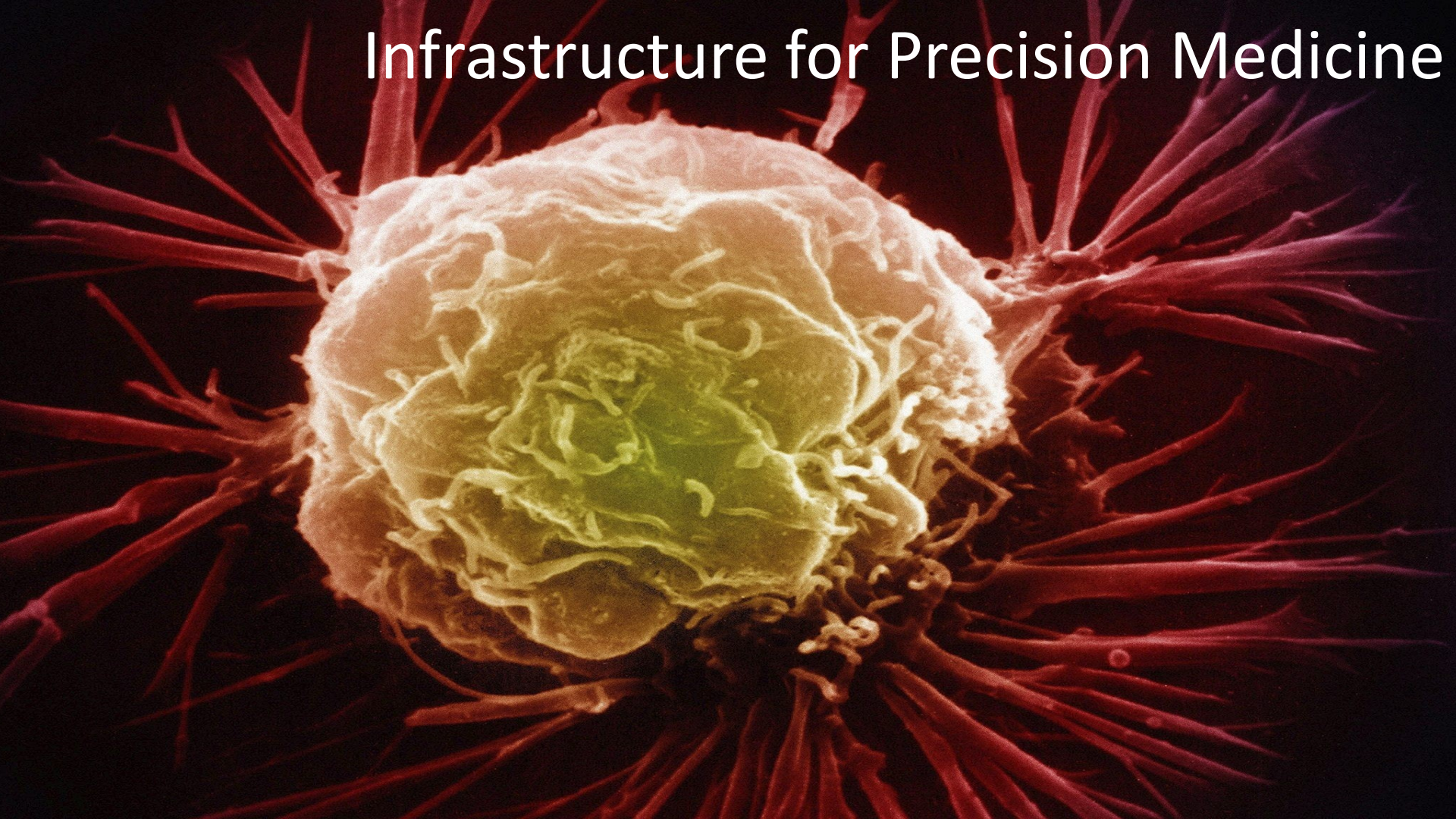


- Harmonization and storage for the Nations Cancer Genomic Data GDC 1.6PB of cancer genomic data and associated clinical data.
- **Precision Medicine Enabled By Precision Networking**

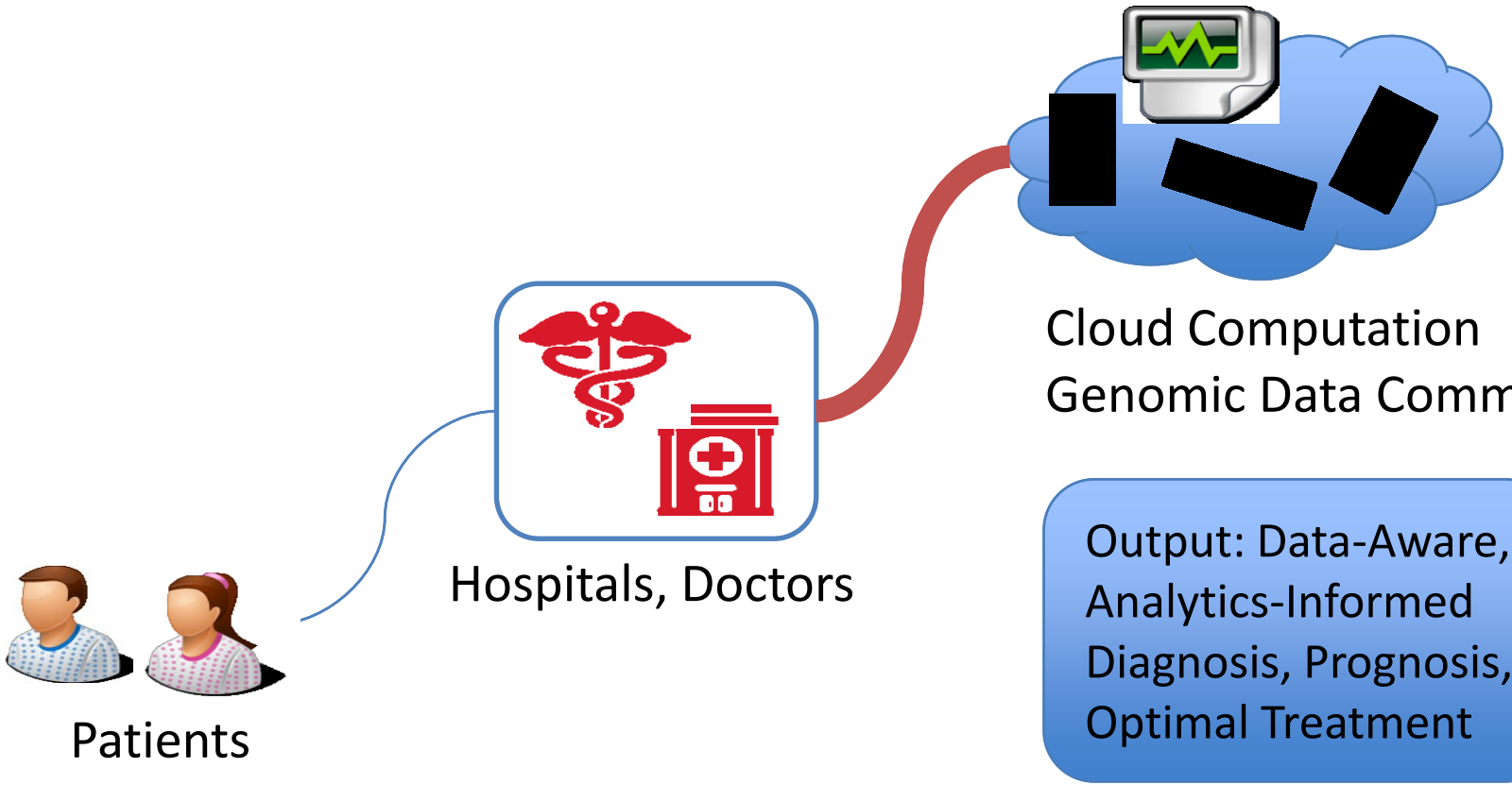
Bionimbus Protected Data Cloud

- Petabyte-scale, secure compliant biomedical cloud that interoperates with dbGaP controlled access data at NIH.

Infrastructure for Precision Medicine



Future Vision: A Nationwide Virtual Comprehensive Cancer Center



Opportunity: Close Integration of Research Workflows and Foundation Networks

- Opportunity: Using GENI To Develop Innovative Techniques for Extremely Close Integration of Research WorkFlows and Dynamic Programmable Network Resources, Enabling Precision Networking
- Network Foundation Architecture: GENI + Innovative Customized Software Defined Networking Exchange (SDX)
- For This Demonstration: Specifically To Meet The Requirements of Bioinformatic Workflows

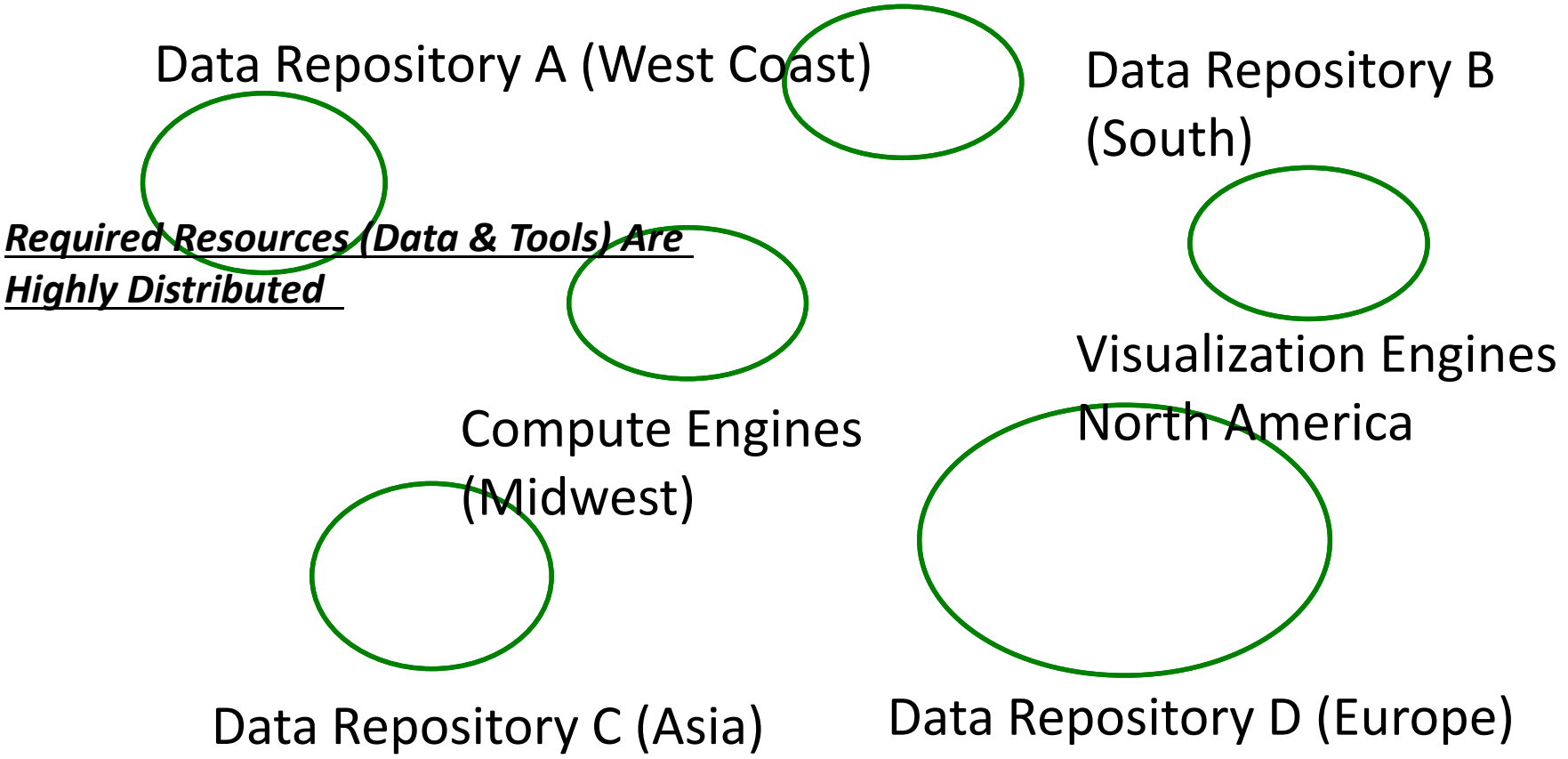
GENI Network Programmability Is Key

GENI Programmability

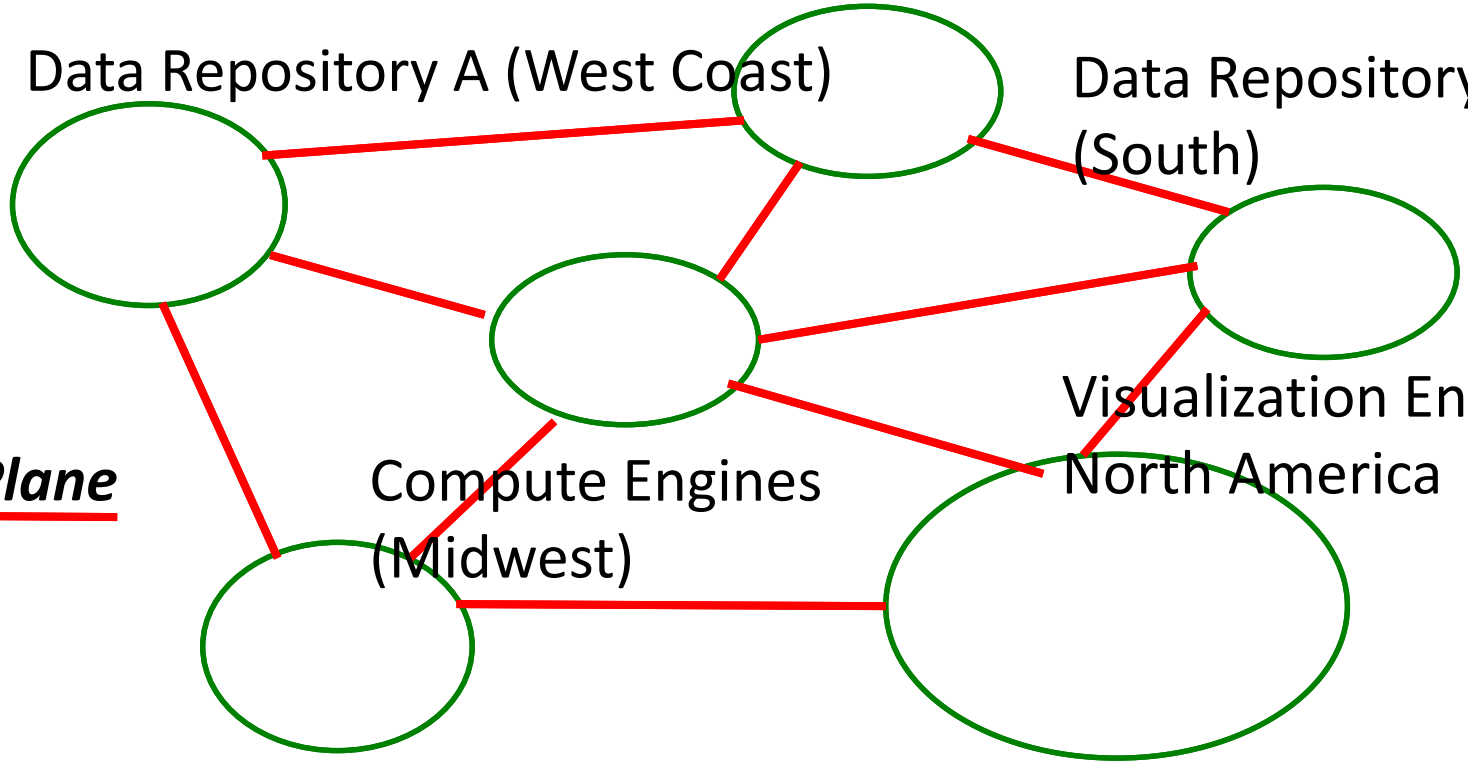
- GENI Provides A Platform for Building the Required Precision Communication Services, Networks and Exchanges (SDXs)
- GENI OpenFlow Network
  - National Overlay Infrastructure Comprised of Shared VLANs Interconnected With OpenFlow Switches
  - FOAM/FlowVisor Enabling Sliced OF Switches (e.g., via Subnet, VLAN, Tunnel, etc)
- Discoverable, Integratable, Configurable, Programmable, Virtual Devices: Click Routers, OVS Switches, Mobile Devices, Instrumentation, and Other Resources
- Dynamic Edge Process Topology Design and Implementation

Precision Networks for Precision Medicine

Biomedical Data Commons

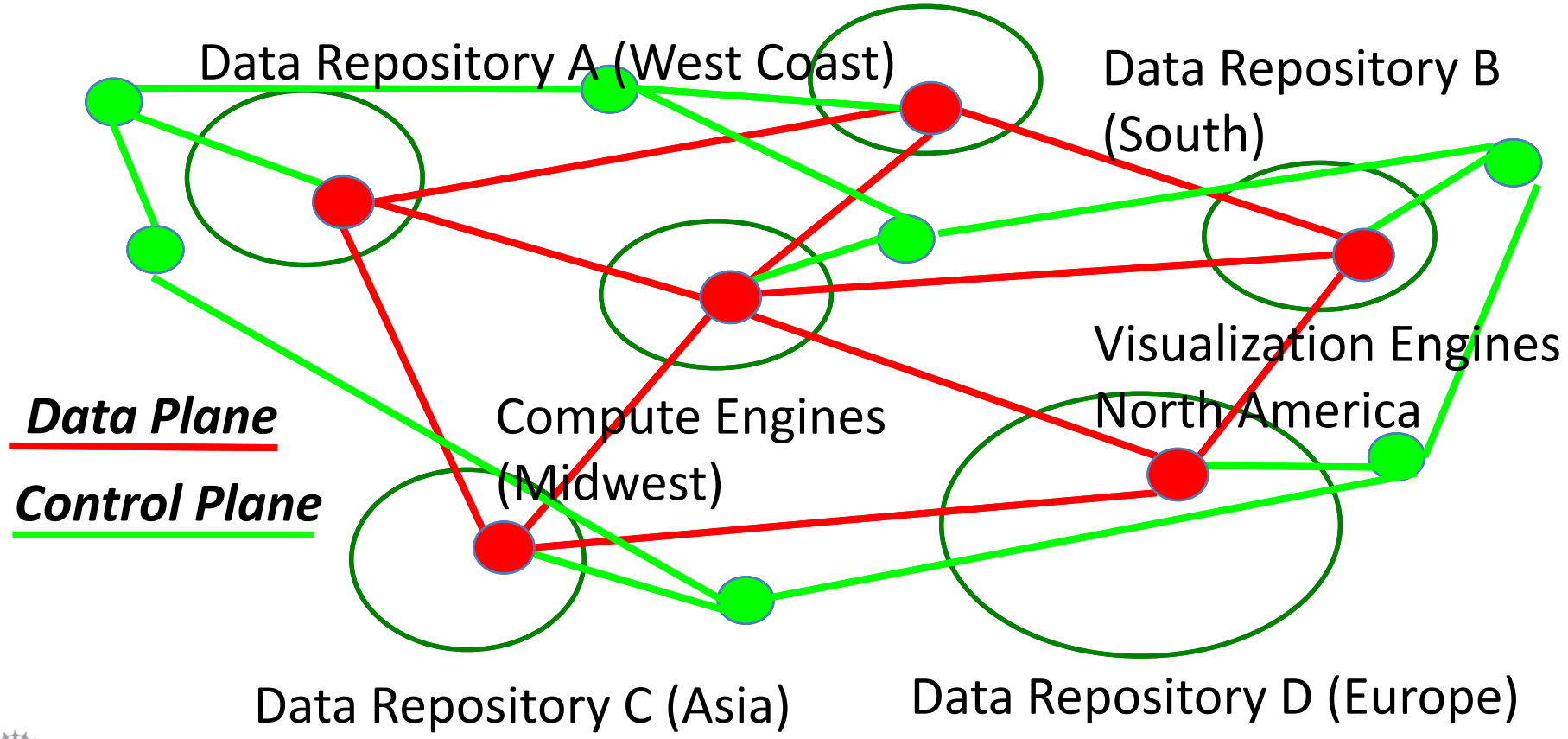


Biomedical Data Commons:
Flow Orchestration: Data Plane

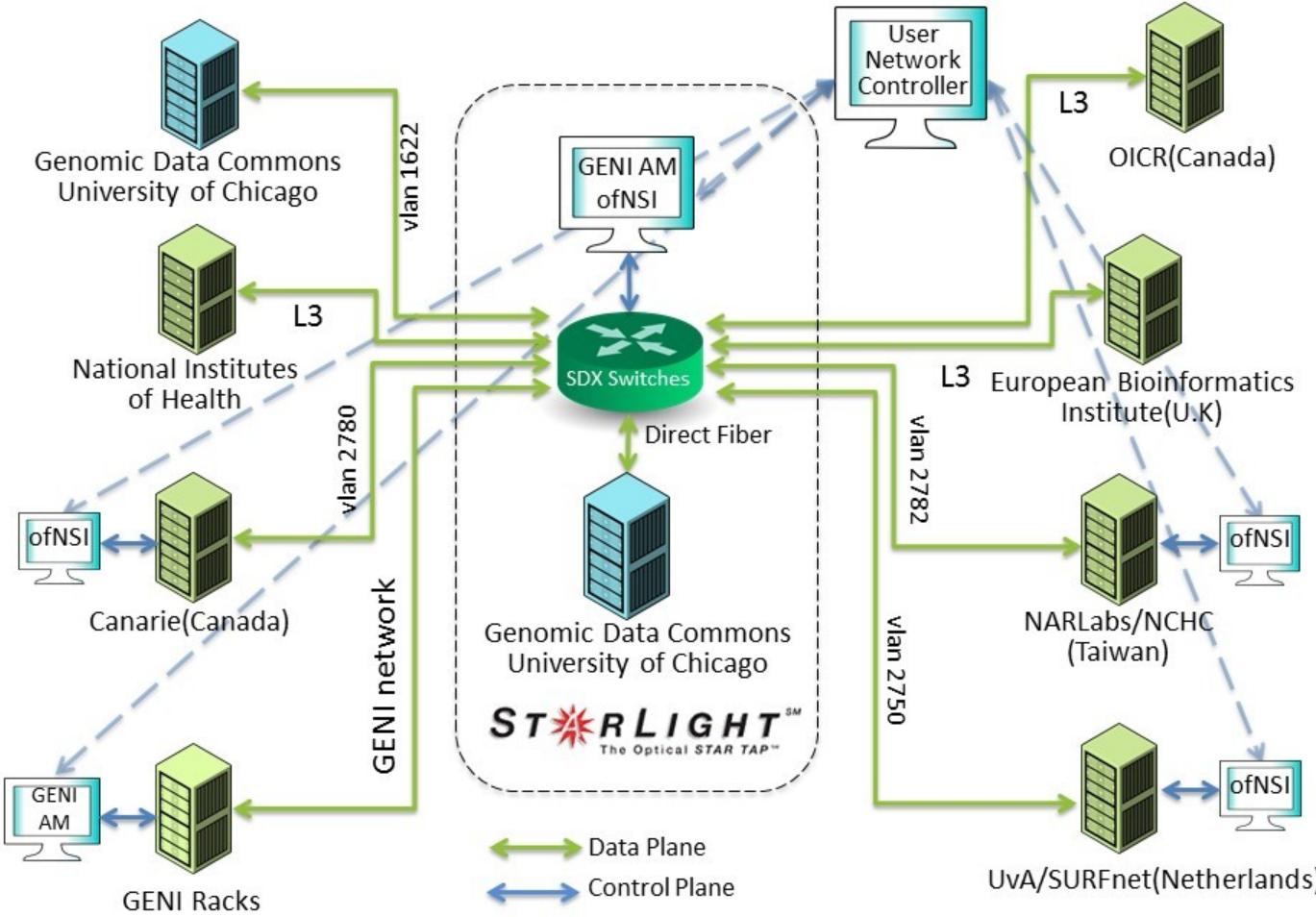


Data Plane

Biomedical Data Commons:
Flow Orchestration: Control Plane + Data Plane



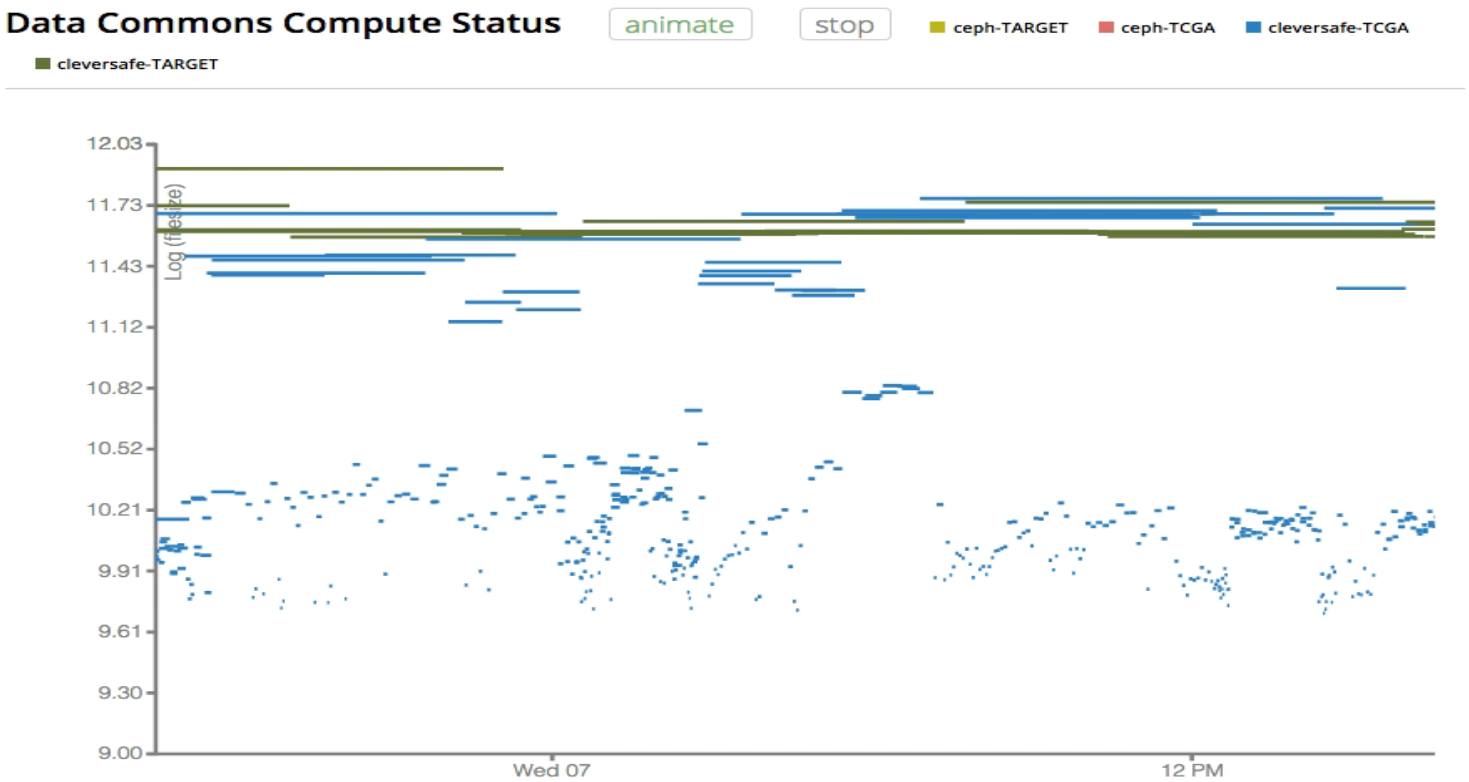
GEC22 Bioinformatics SDXs Demo Network



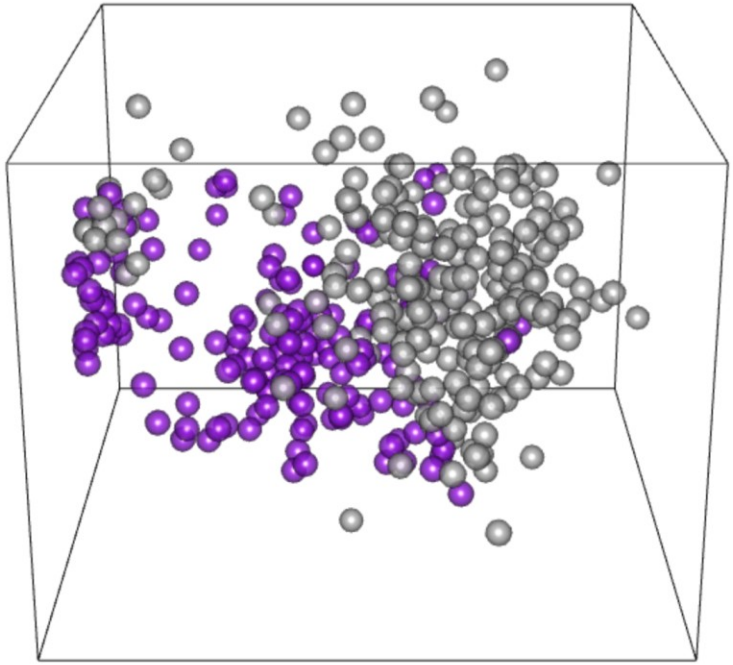
Today's Demonstration

- A) Dynamically Moving Core Data Files Among Multiple Sites Around the World Via StarLight SDX
- B) Moving RNA-seq Data Files From NCI (Bethesda, MD) and EBI (Hinxton, UK) Through SDX Switch/Routers to The University of Chicago.
  - Analysis By Comparison To Known Data Correlated To Drug Response.
  - Determine Possible Actionable Therapeutic Options.
  - Return Viable Treatment Options To the Originating Site.

Genomic Data Commons Data Transfer



Gene Expression Clustering of Lung Cancers

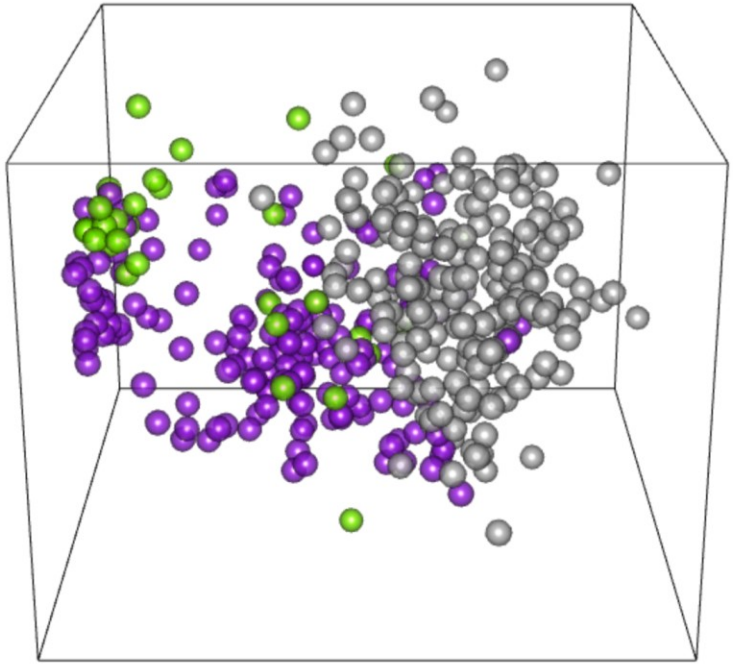


Color By:

Original Diagnosis

- Lung squamous cell carcinoma (LUSC)
- Lung adenocarcinoma (LUAD)

Gene Expression Clustering of Lung Cancers

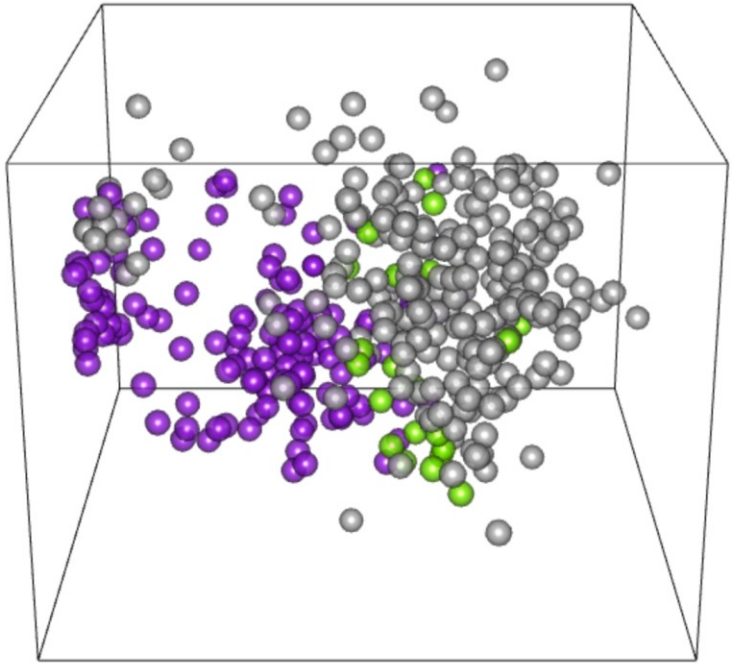


Color By:

Potential Misdiagnosis os LUSC

- Lung squamous cell carcinoma (LUSC)
- Lung adenocarcinoma (LUAD)

Gene Expression Clustering of Lung Cancers

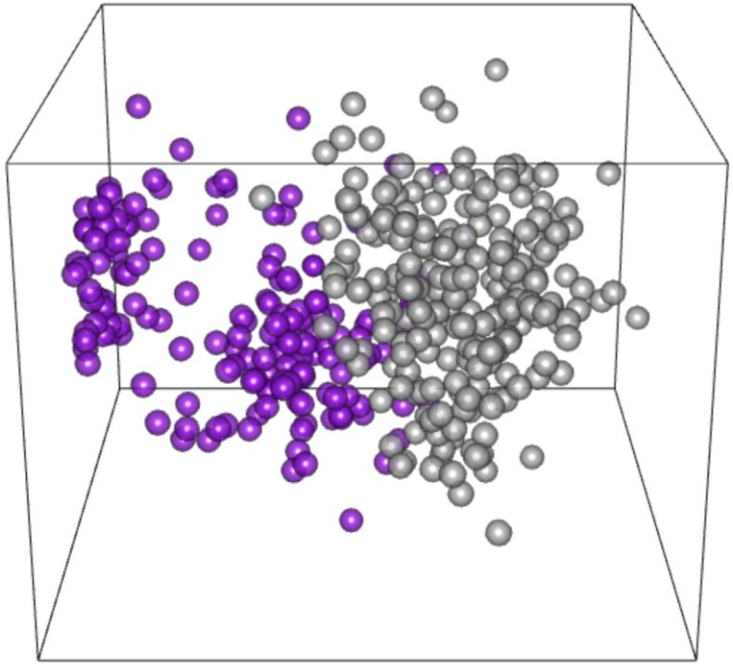


Color By:

Potential Misdiagnosis of LUAD

- Lung squamous cell carcinoma (LUSC)
- Lung adenocarcinoma (LUAD)

Gene Expression Clustering of Lung Cancers



Color By:

By Expression

Lung squamous cell carcinoma (LUSC)
Lung adenocarcinoma (LUAD)

Results

- Precision medicine requires data commons that scale to hundreds of petabytes scale, with programmable networks and data peering to support data sharing.
- Speed discovery and support analytics-driven healthcare to recommend treatment.
- Large Scale Data Analysis and Dynamic Pipelines For Workflows Are Essential For Determining Optimal Results.

Summary and Future

- What you saw: An innovative approach to advanced knowledge discovery and medical treatment***:*** **Precision medicine being supported by precision networking**
- Why GENI/US Ignite is important: Precision mapping of communication services to BI workflow requirements across the world using advanced analytics, the Genomics Data Commons & a programmable dynamic SDX
- What happens looking forward, for the application and its integration with GENI:
  - A) Further development/refinement of basic capabilities
  - B) Transition to ***actual production services***
  - **C) The Genomics Data Commons and Bionimbus Protected Data Cloud is Being Developed As a Key Production Knowledge Discovery/Transformational Medical Treatment Facility**

Using GENI To Invent the Future…

Thank You!

usignite geni