

Genomic Studies at Inova



Inova Translational Medicine Institute
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Organization

- Background
- Data management
- Challenges \ Lessons Learned



Background: The Healthcare System

Six hospital + ambulatory healthcare system

- Largest healthcare system in Northern VA
- Two million patient visits/year
- 20,000 deliveries/year

Main hospital

- 1,000 beds including Children's hospital
- 10,000 deliveries/year



Background: The Research Institute

- Started: 2010
- Overall goal: research on integration of genomic information into the practice of medicine
- Staffing: 1/3 clinical, 1/3 bioinformatic/IT, 1/3 laboratory



Background: ITMI Studies

Themes

- Trio-based WGS
- Other 'omics
- Comprehensive clinical data
- Integrated Laboratory
- Unified database



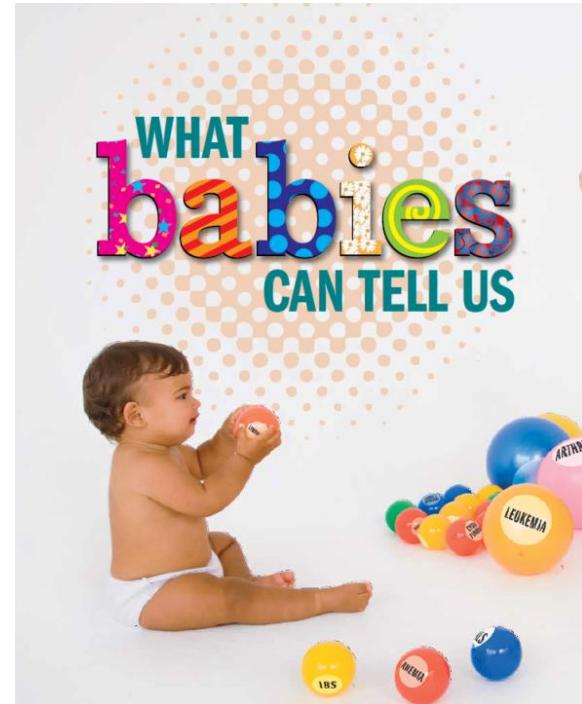
Preterm Birth Study (2011)

- Molecular associations with preterm birth
- ~500 PTB trios, ~500 FT trios
- WGS + 'omics + clinical data
- Specimens: blood, saliva, cord blood, placenta



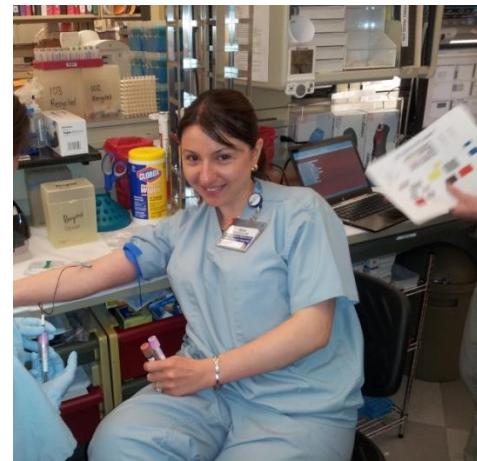
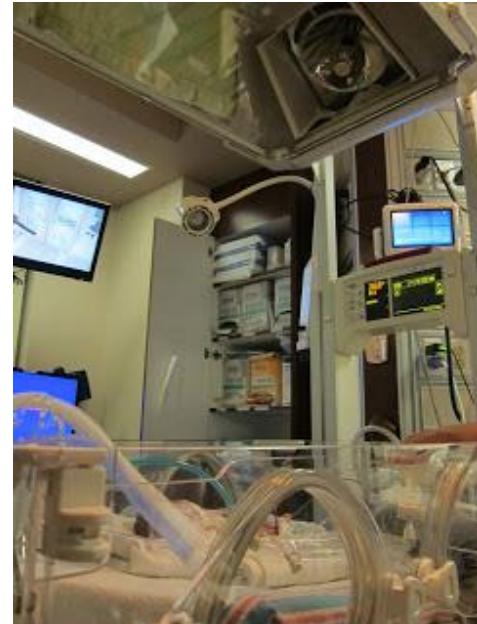
Longitudinal Study (2012)

- WGS + 'omics + clinical data on 5,000 → 10,000 trio-based families
- Longitudinal study (≥ 18 yrs)
- Blood, saliva, urine, cord blood, placenta
- DNA, RNA, protein, epigenetic + clinical data



Congenital Disorders Study (2012)

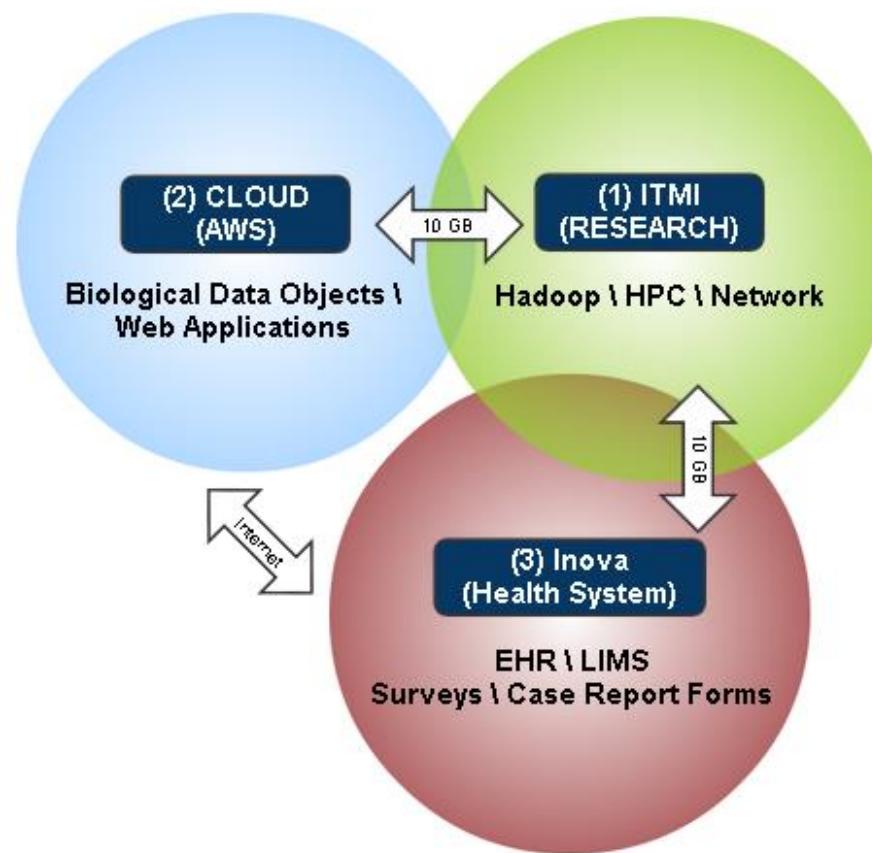
- Mostly NICU-based
- Any other patient with a “congenital/genetic” disorder
- ~2-3 families/week
- Trio-based WGS, etc.



Data Management

Infrastructure: Hybrid Cloud

Cloud



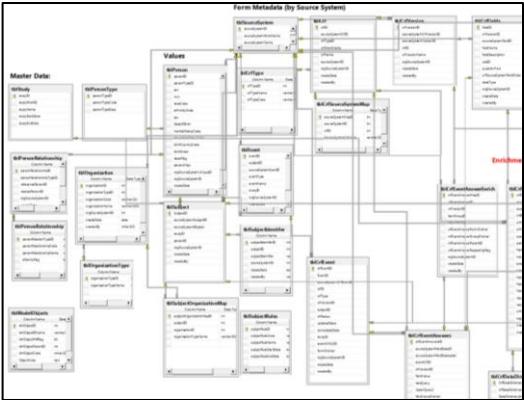
On-Premises

Database

Yes, a Hybrid

Relational Databases

Inova Infrastructure



Object Store

AWS

Storage & Content Delivery



S3
Scalable Storage in the Cloud



Storage Gateway
Integrates On-Premises IT Environments with Cloud Storage



Glacier
Archive Storage in the Cloud

NoSQL \ Hadoop
Value pair \ Graph
Research Network

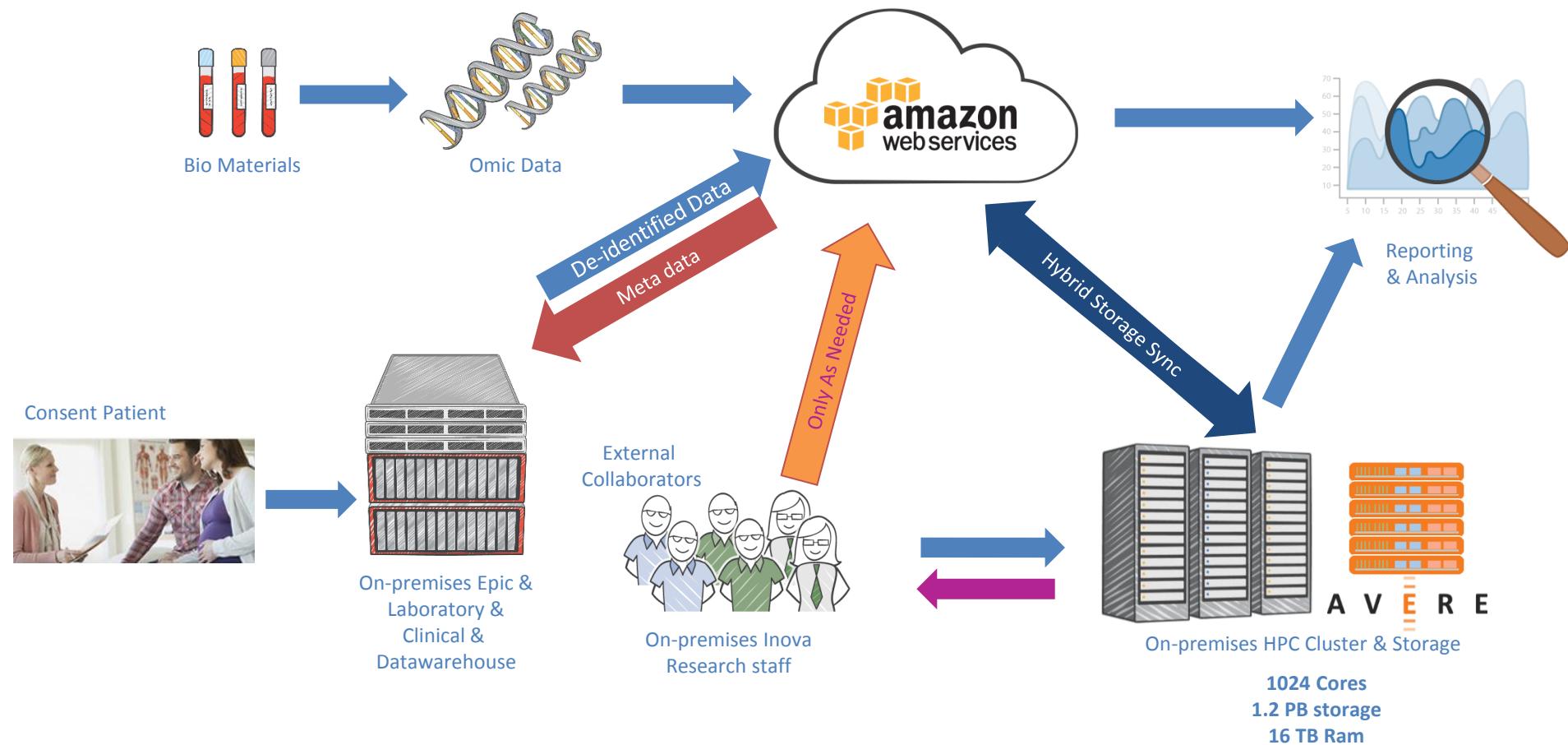
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"fields": [
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  {"name": "clientProtocol", "type": ["null", "string"]},
  {"name": "serverHash", "type": "MD5"},
  {"name": "meta", "type": ["null", {"type": "map", "values": "bytes"}]}
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]
  
```

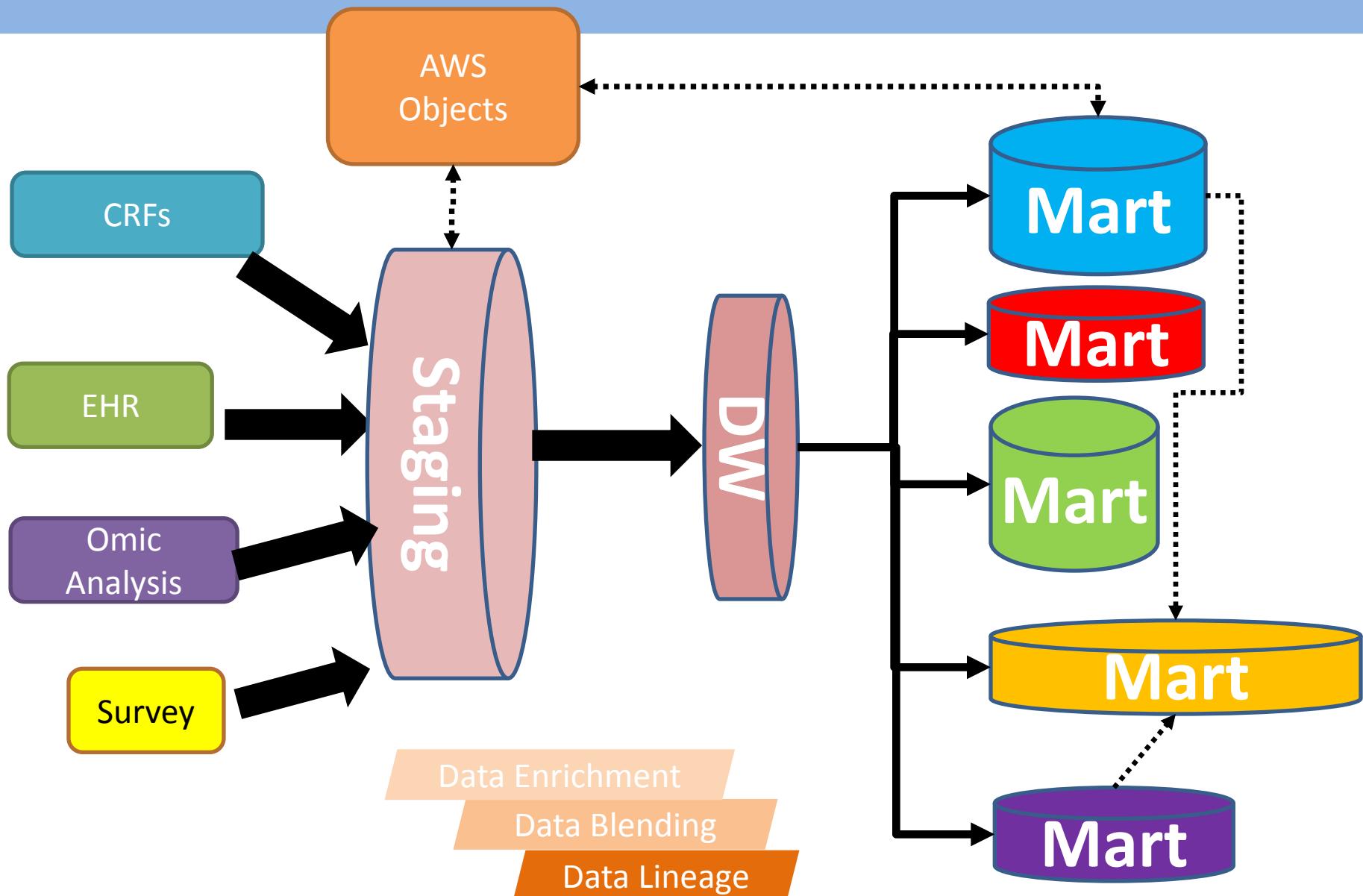
ITMI Data Collection



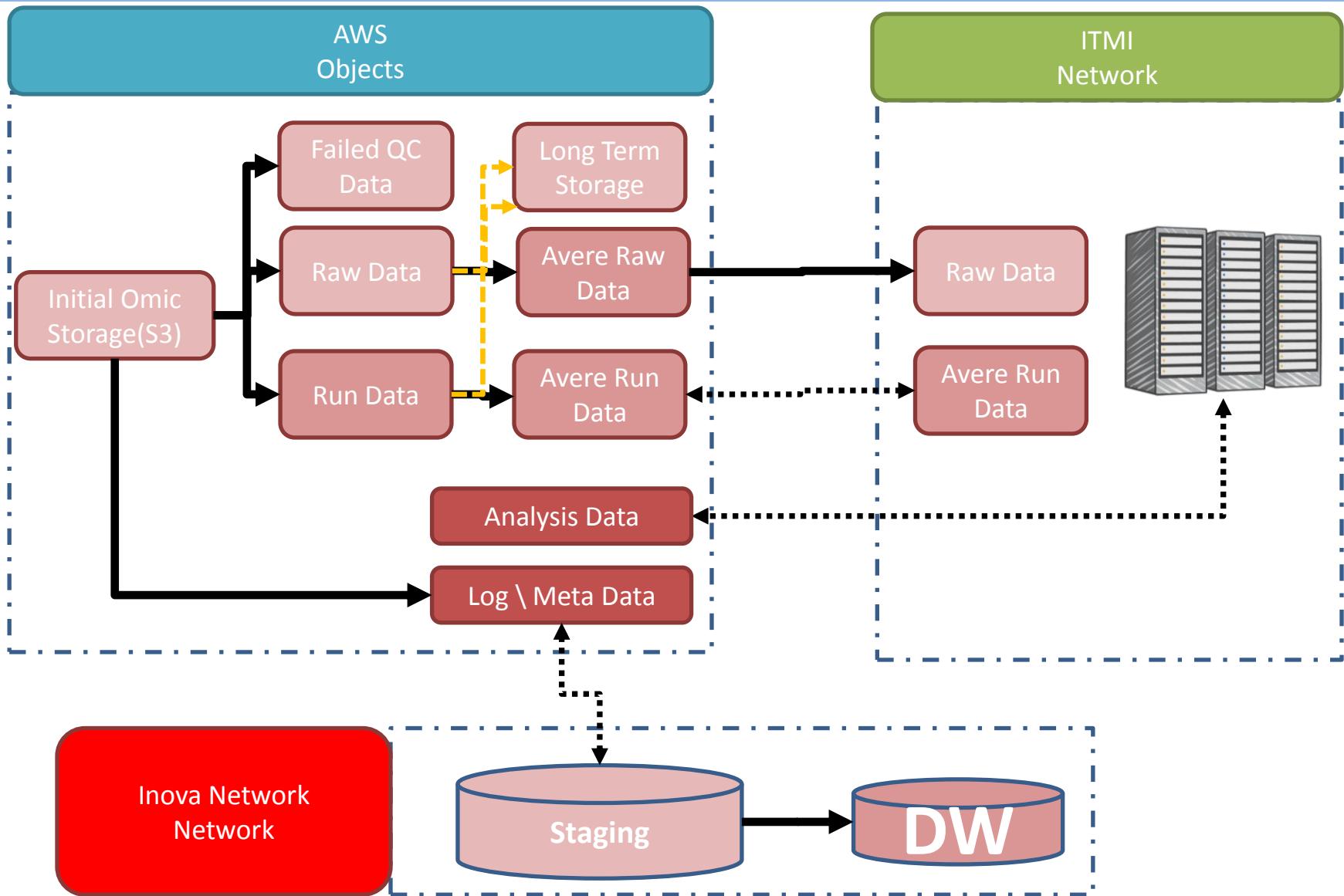
The Stats

- **Participants**
 - **9,750** participants (enrolled), **>110** Different Countries of Birth
- **Files and data**
 - Manage **3+** Petabytes of storage capacity (between cloud and on-premise)
 - **~10,000,000** million files
 - File Sizes Range from Kilobytes to **100+ GB** per file
- **Specimens**
 - **>450,000** specimens
- **Whole Genomes Sequences**
 - **7,300 +**
 - **~36,500,000,000 variants!**
 - *Also have epigenetic data*
- **Clinical**
 - **55,000+** Patient Diagnosis (Longitudinal)
 - **110,000+** Surveys and Case Report Forms
- **Labs Results**
 - **2,000,000+** discrete lab results
 - **2,200,000+** discrete variables from Case Report Form and Surveys

General Data Process

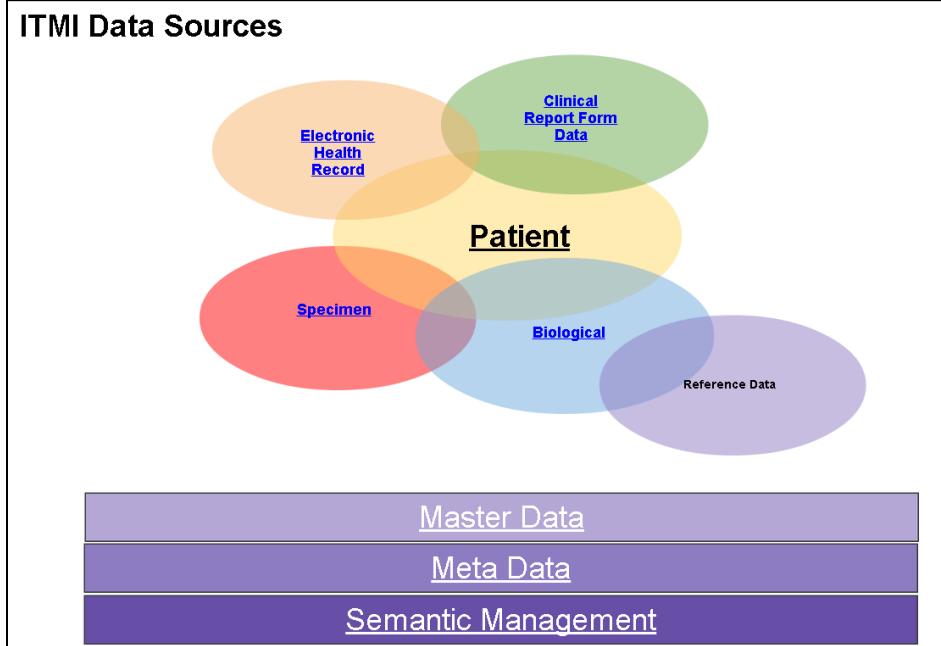


Omic Data Process



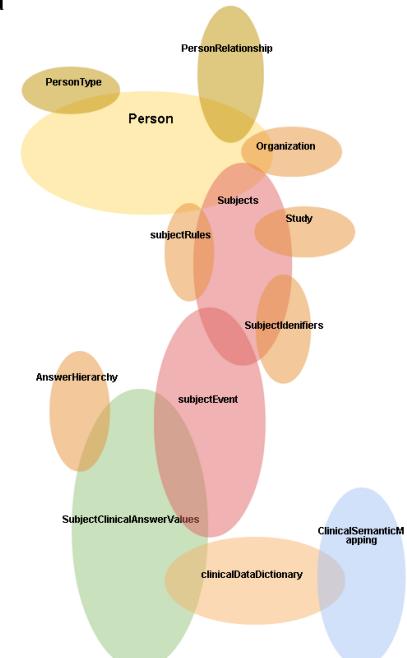
Data Model

(1) Logically Model

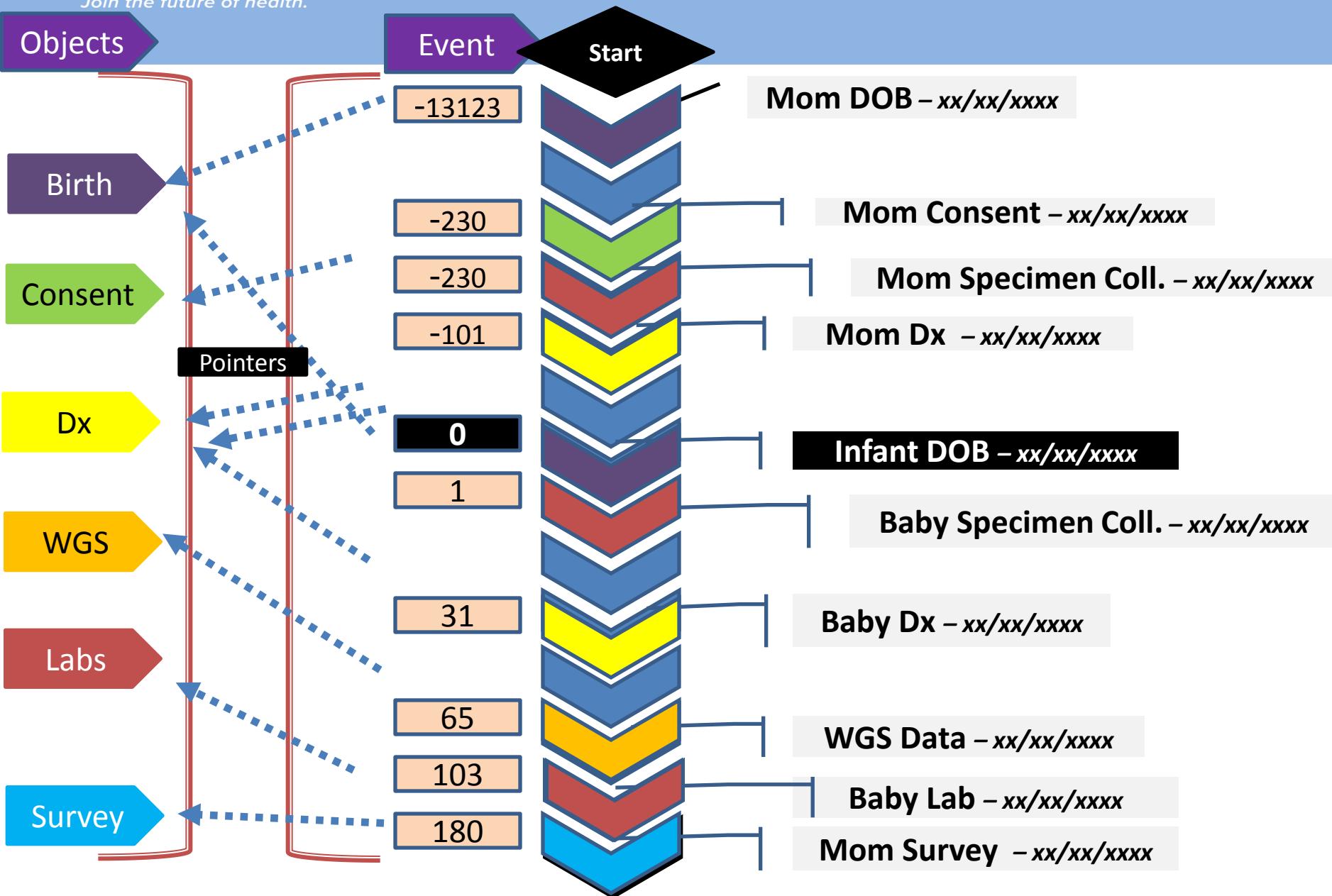


(2) Break down each

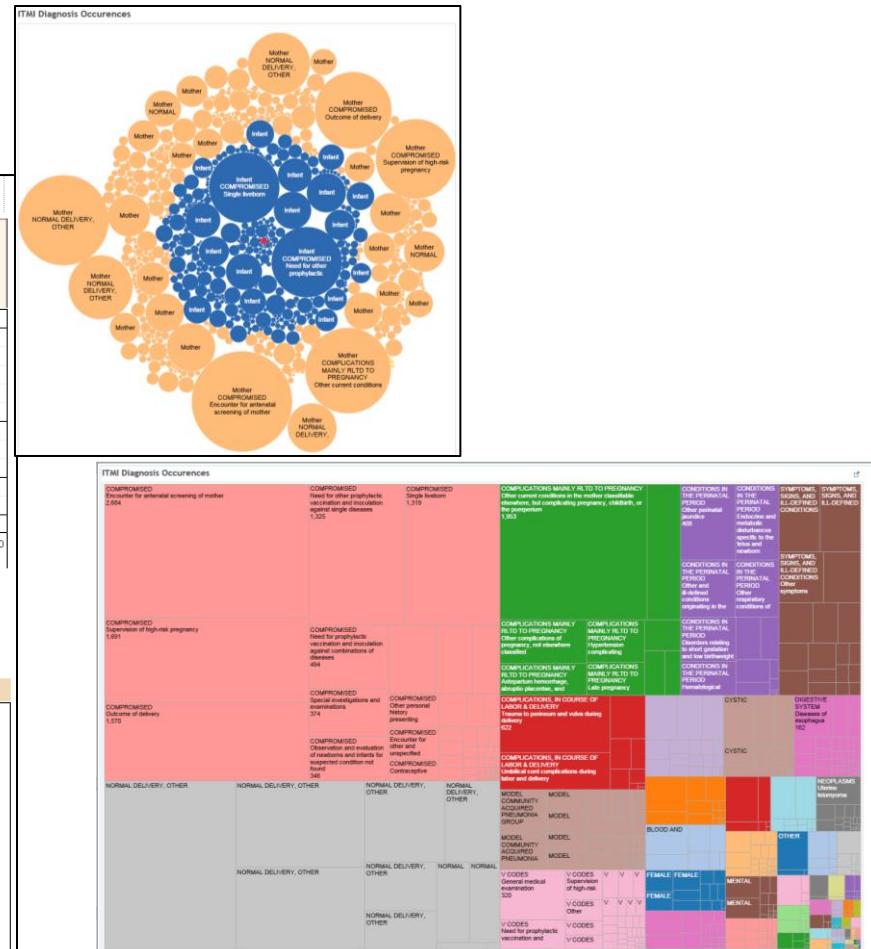
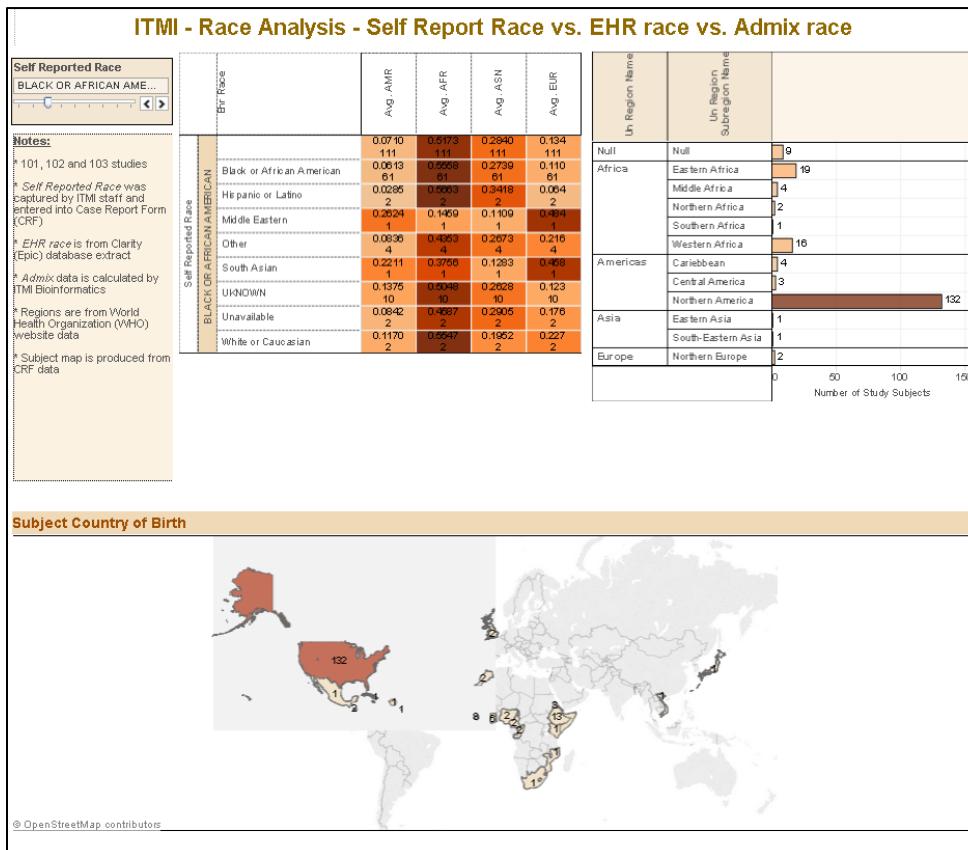
Logical Model
Report Form Data



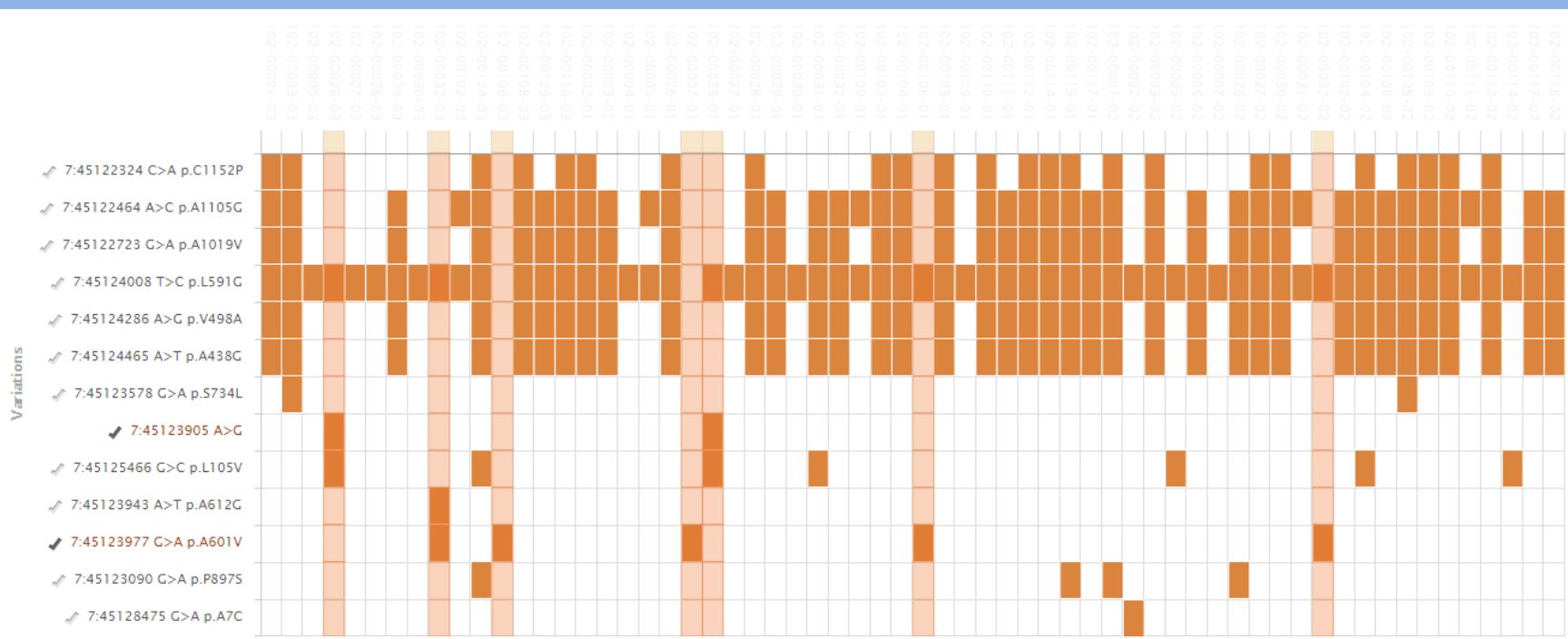
LONGITUDINAL DATA



Visualization



Web Portal


 Select All

i

DISPLAY SETTINGS

 MISSENSE VARIANT

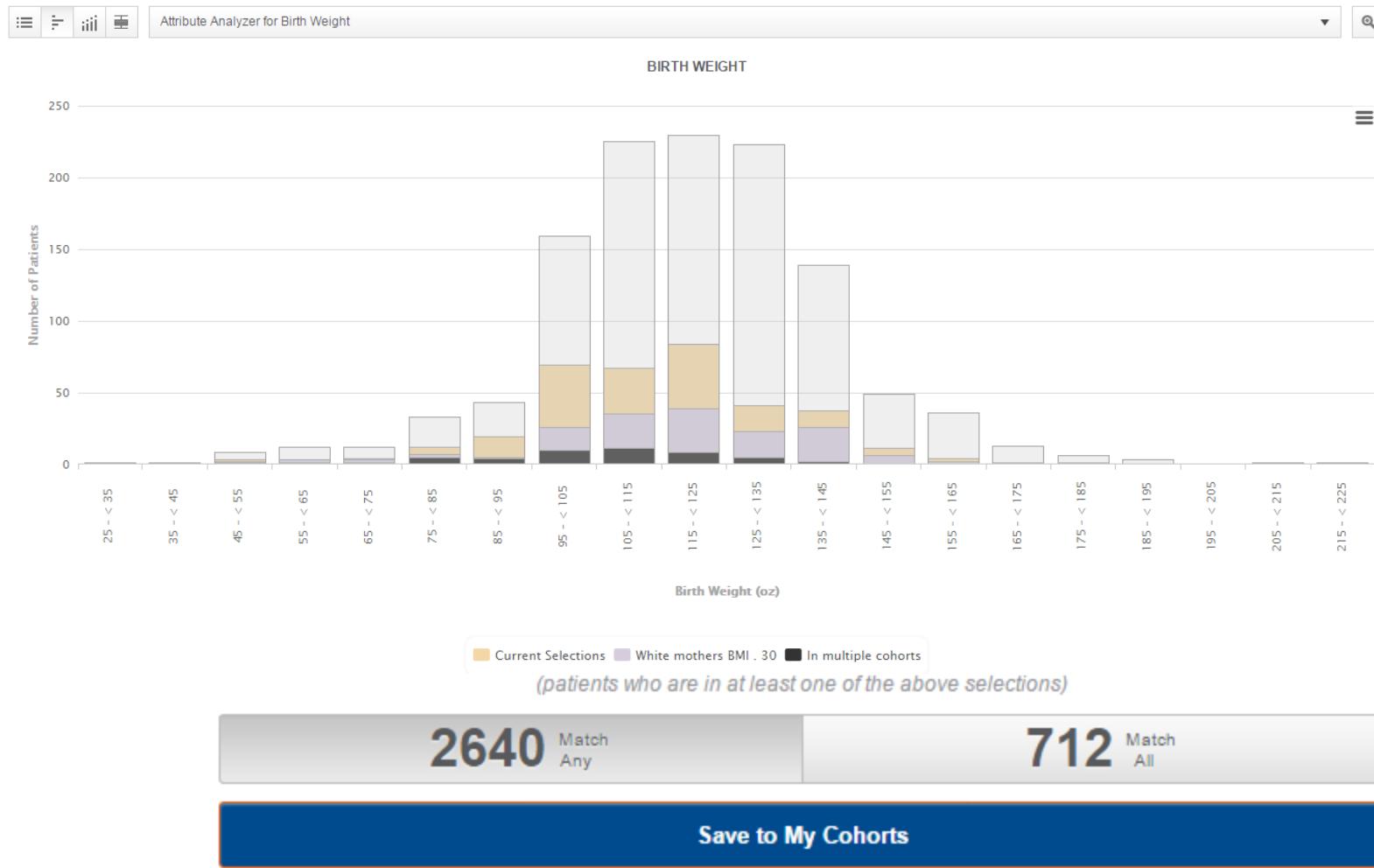
62

62 patients have mutations out of 74 total.

 Only show patients with mutations

Start Tutorial

Web Portal



III. Challenges \ Lessons Learned

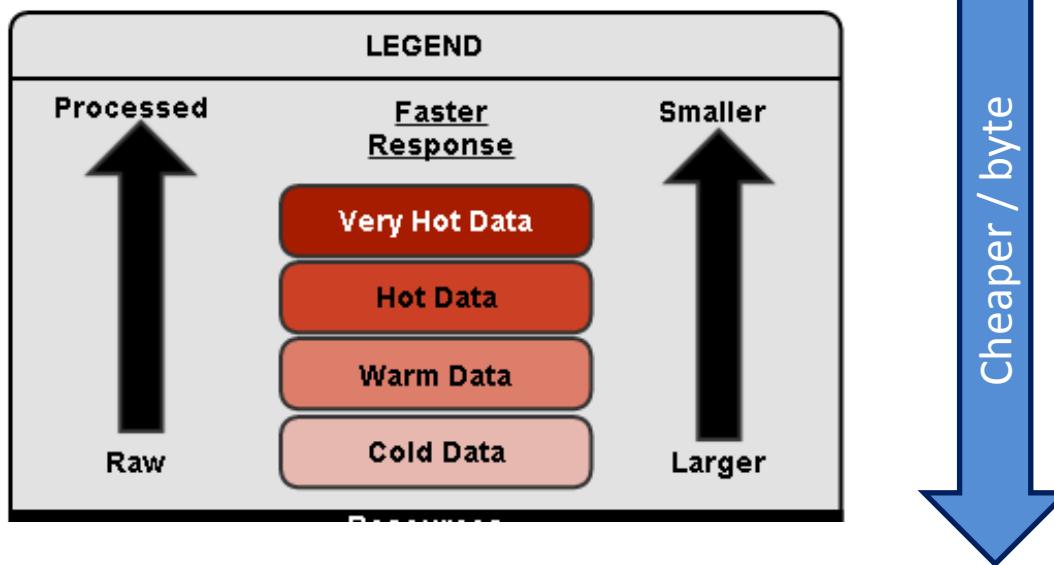


Challenges

- Scalability
 - Storage
 - Compute
- Data Movement
- IT Standards
 - Data
 - Programming

Lessons Learned

- Build infrastructure around data
 - Network is bottleneck
 - Hidden costs
- Manage Data Tiers



Lessons Learned

- Spend time to model
 - Build strong metadata layer
 - Make it understandable to your team
 - Use best practices
- IT Partners that understand the business

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