



Icahn School  
of Medicine at  
**Mount  
Sinai**

# Mount Sinai Data Warehouse Town Hall

May 15, 2025

# The MSDW Team



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Database Administrator



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# Agenda

## 1 MSDW Operations

## 2 MSDW Major Accomplishments

- Integration of Somatic Genomic Results into Leaf
- Digital Pathology Slide Diagnoses

## 3 Epic for Research

## 4 MSDW Roadmap May 2025 – November 2025

# MSDW Operations

# MSDW Data Sets Delivered

- From October 2024 to April 2025, the MSDW team closed 87 data requests
- There has been a significant decrease in the data set delivery time from October 2024 to April 2025 due to increased operational efficiency
  - Increased operational efficiency by implementing KPIs and SLAs

## Top Department Requesting MSDW Data Sets

Apr 2024 – Sept 2024

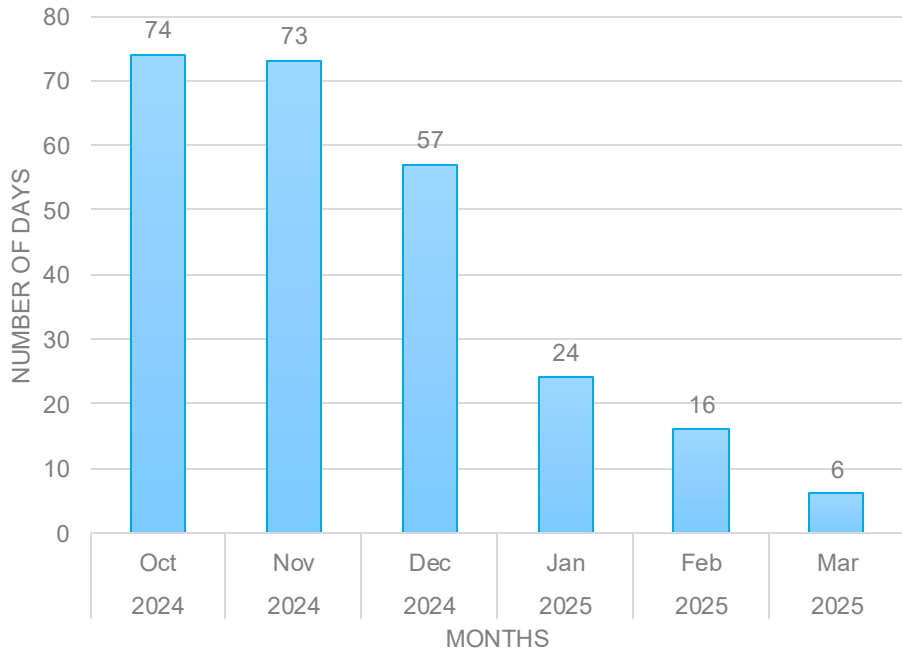
Department	Tickets (Number)
Population Health	15
Genetics and Genomics	10
Medicine	9
Medical Oncology	8
Psychiatry	7

Oct 2024 – Apr 2025

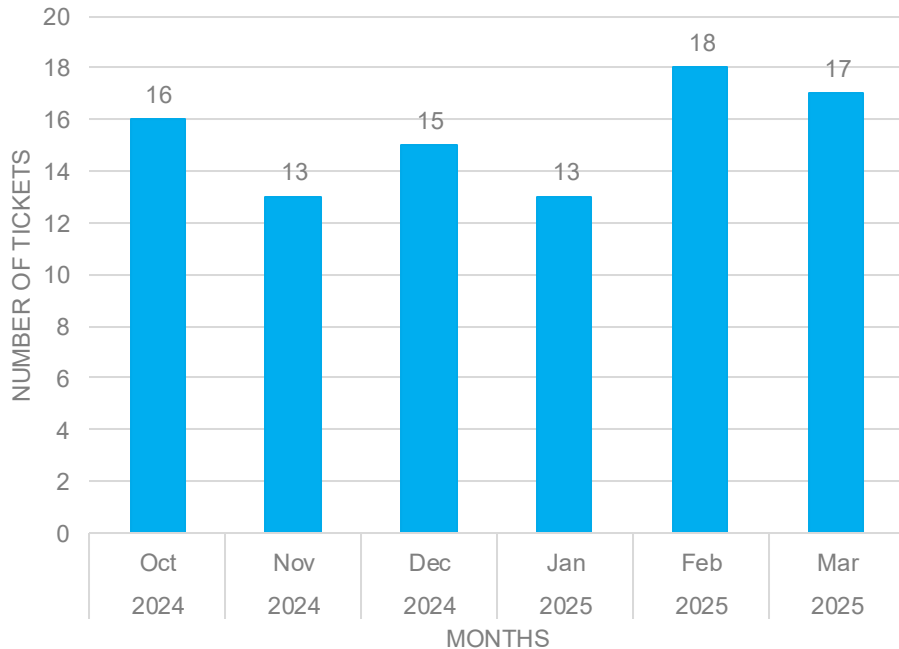
Department	Tickets (Number)
Medicine	22
Genetics and Genomics	16
Psychiatry	15
Population Health	14
Surgery	12

# Custom Data Set Average Days to Resolution

Average Days to Resolution



Number of Tickets Closed

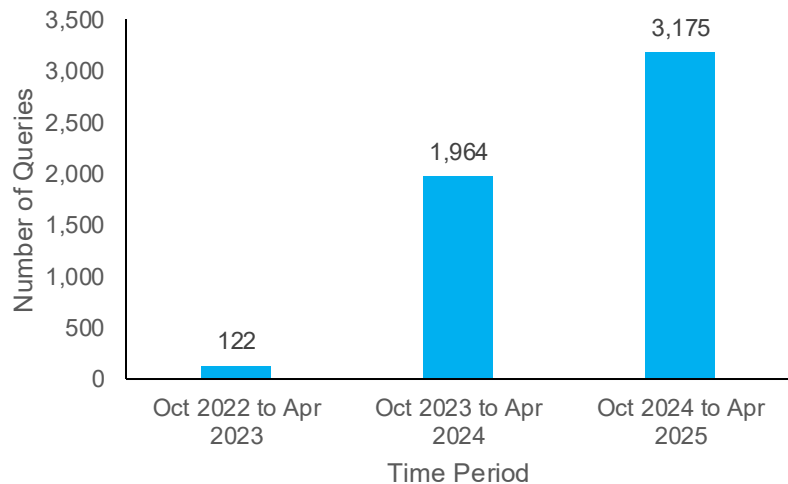


# Leaf Utilization: Patient Cohorts & Custom Search Facets

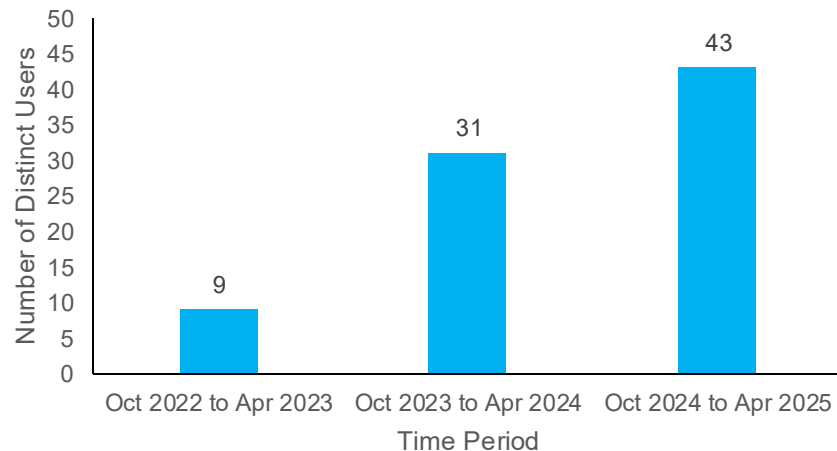
Patient Cohort / Feature	Queries Executed (N)	
	April 2024 – Oct 2024	Nov 2024 - Apr 2025
Cancer Staging from Cancer Registry	196	153
BioMe BioBank	102	102
Imaging Modality/ Body Part from IRW 2.0	NA	80
Digitized Pathology Slides Cohort	5	29
BioMe BioBank Global Diversity Array (Sema4)	6	16
BioMe BioBank Whole Exome Sequencing (Regeneron)	0	5
Imaging Research Warehouse 1.0	0	4
Cancer Patient Cohort	2	4
Cancer Institute Biorepository	14	3
Imaging Research Warehouse 2.0	9	1
BioMe BioBank Global Screening Array (Regeneron)	0	0
<b>Total Leaf Queries</b>	<b>797</b>	<b>1006</b>

# TriNetX Utilization Continues to Increase

Number of Queries Run between  
October 2024 & April 2025 Compared  
to Previous Years



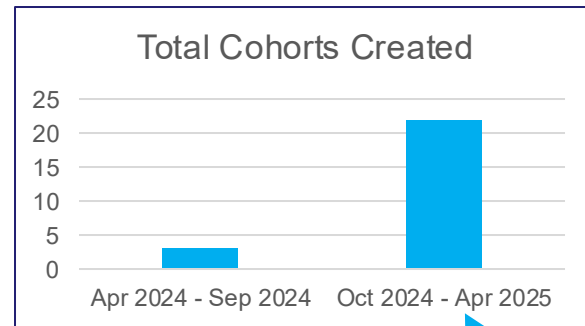
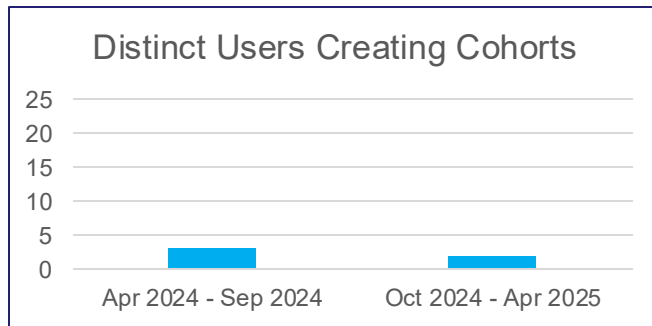
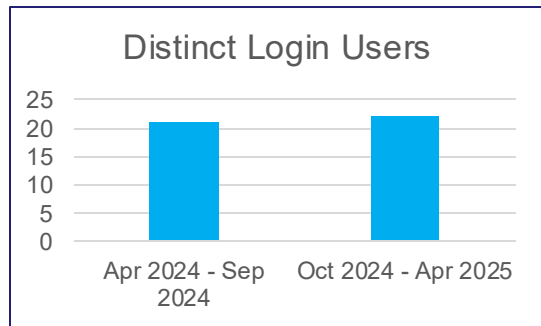
Number of Unique Users between  
October 2024 & April 2025 Compared  
to Previous Years



**There has been a substantial increase in both users and queries run since the previous year's report**



# ATLAS Utilization: April 2024 – April 2025



- Utilization of ATLAS remains low with few users who are logging in creating and saving cohorts

A single user created 21 cohorts in October 2024. This user has not logged in again since October 2024

# Outreach to MSDW Users: April 2024 – April 2025

Event	Participants
Epic for Research Training	200
TriNetX Training Session	30
Leaf and ATLAS Training Session	36
Digital pathology training session	62
Leveraging Electronic Health Record (EHR) Data for Data Analysis	66
Digital Concierge	410
<b>Total</b>	<b>623</b>

# MSDW Major Accomplishments

# Obtaining Somatic Genomic Results from External Vendors

## ► Project objectives

- To link the phenotypic and somatic genomic data on Minerva facilitating the use of somatic genomic data for analytics, clinical research and clinical operations
- To make raw and structured somatic genomic results from external vendors available to the Mount Sinai research and clinical community

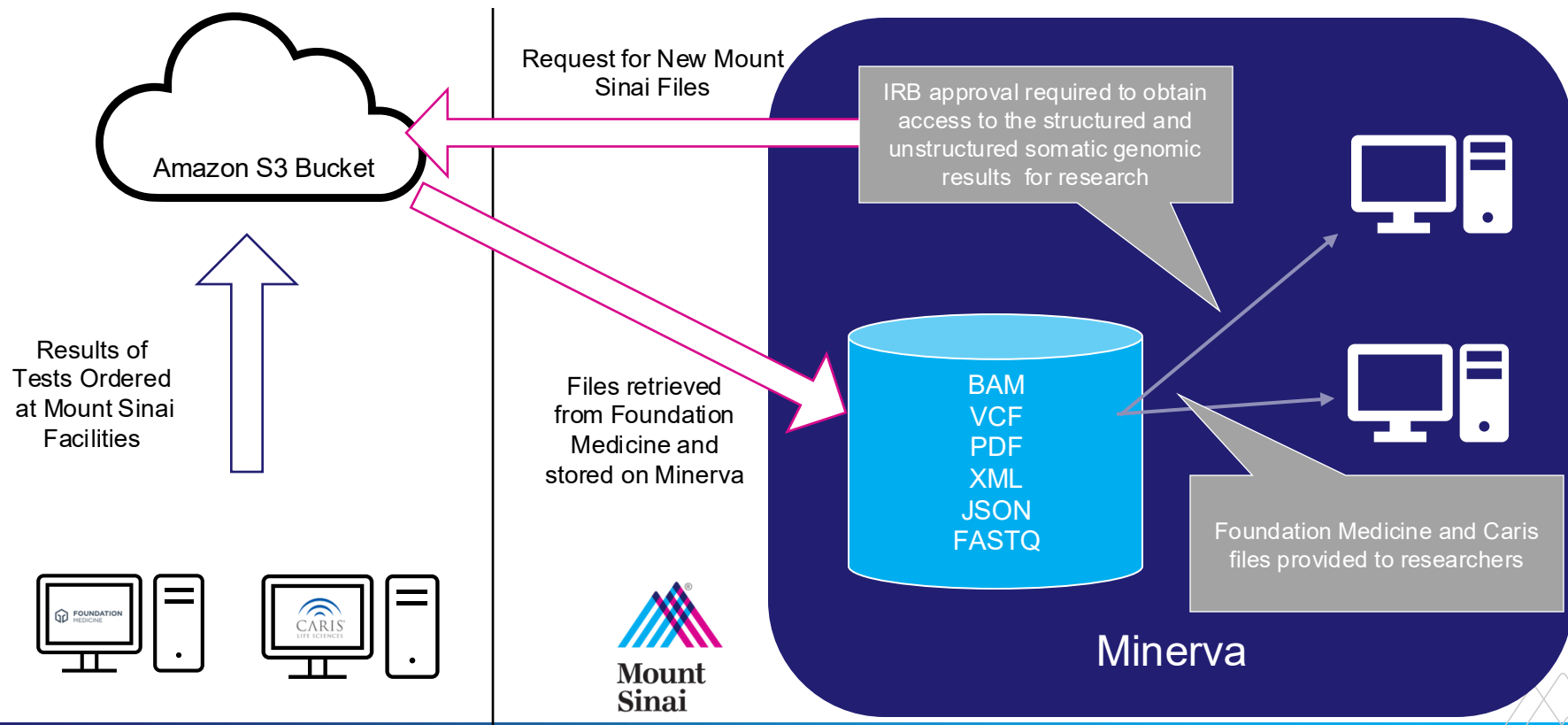
## ► Collaboration with Mount Sinai Innovation Partners (MSIP)

- MSIP ensuring contracts address Mount Sinai's best interest for use of somatic genomic data

## ► Genomic results returned to Minerva

- Results include both structured and raw genomic data
- File formats received include BAM, FASTQ, VCF, PDF, XML, JSON and CSV
  - File types available vary by vendor

# Data Flow from Somatic Genomic Testing Vendors to Mount Sinai



# Search Select Genetic Mutations in Leaf

The screenshot shows the Leaf Genomics web application interface. The top navigation bar includes the Leaf logo, a search bar, and a 'Save Query' button. The main content area displays a list of genetic mutations under the 'Somatic Genetic Mutations' category. A search filter is applied, showing results for 'EGFR' mutations. A magnifying glass highlights the 'EGFR' section, showing a list of specific mutations and their patient counts.

Search Results for EGFR:

Mutation	Count
EGFR	317
EGFR (exon 19 Deletion)	60
EGFR (L858R)	48
EGFR (Other/ Unspecified Mutation)	202
EGFR (T790M)	7

# Pathogenic Gene Mutations Exposed in Leaf

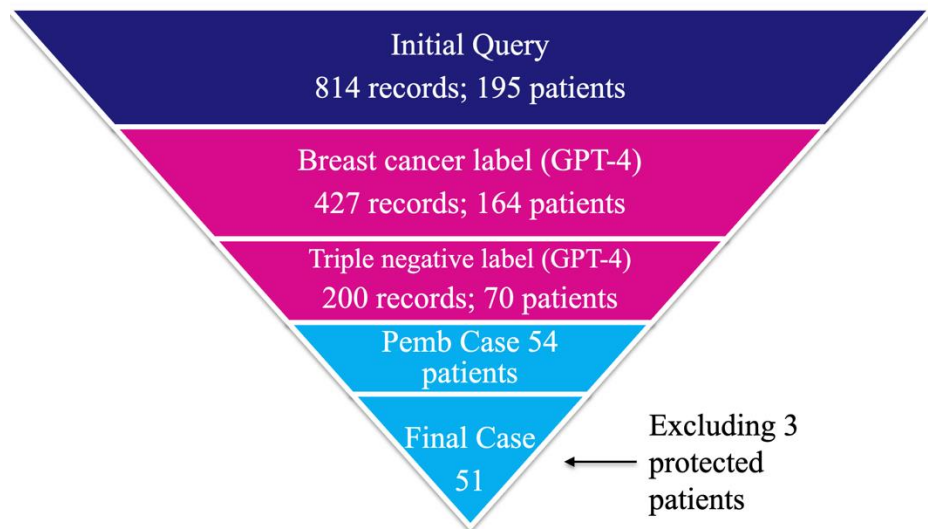
GENE	MUTATION DETAILS	GENE	MUTATION DETAILS	GENE	MUTATION DETAILS
ALK	<i>ELM4-ALK fusion</i>	DDR1		MSI-H	
APC		DDR2		MTOR	
ATM		EML4		MYC	<i>Amplification</i>
ATR		EGFR	<i>L858R, T790M, Exon 19 Deletion</i>	NTRK1	<i>NTRK1 fusion</i>
AXN1		ERBB2	<i>Amplification</i>	NTRK2	<i>NTRK2 fusion</i>
AXN2		EZH2		NTRK3	<i>NTRK3 fusion</i>
TCF7L2		FGFR1		PIK3CA	<i>E542K, E545K, H1047R</i>
RNF43		FGFR2	<i>FGFR2 fusion</i>	PTEN	
BRAF	<i>V600E, V600K</i>	FGFR3	<i>S249C</i>	RET	
BRCA1		FLT1		RNF43	
BRCA2		FLT4		ROS1	<i>ROS1 rearrangement</i>
CD274 (PD-L1)	<i>Amplification</i>	IDH1	<i>R132H</i>	TCF7L2	
CDK4		IDH2	<i>R140Q</i>	TP53	
CDK6		KDR		VEGFA	
CTLA4		KRAS	<i>G12C, G12D, G12V, Wild type</i>		
CTNNB1		MET			

# Using AI to Identify Patients for Cohort Inclusion

**Background:** Using Large Language Models (LLM) to enhance cohort identification. Conventional methods (keywords, ICD codes, structured queries) struggle with nuanced clinical criteria.

**Study goal:** Identify **Stage 1,2,3 TNBC** patients who (1) received **neoadjuvant pembrolizumab** (2) underwent **definitive surgery**. **Exclude:** Stage 4 at diagnosis.

**Solution:** Used **GPT-4 via Azure API (HIPAA-compliant)** to process unstructured pathology reports and notes securely.





# Epic for Research

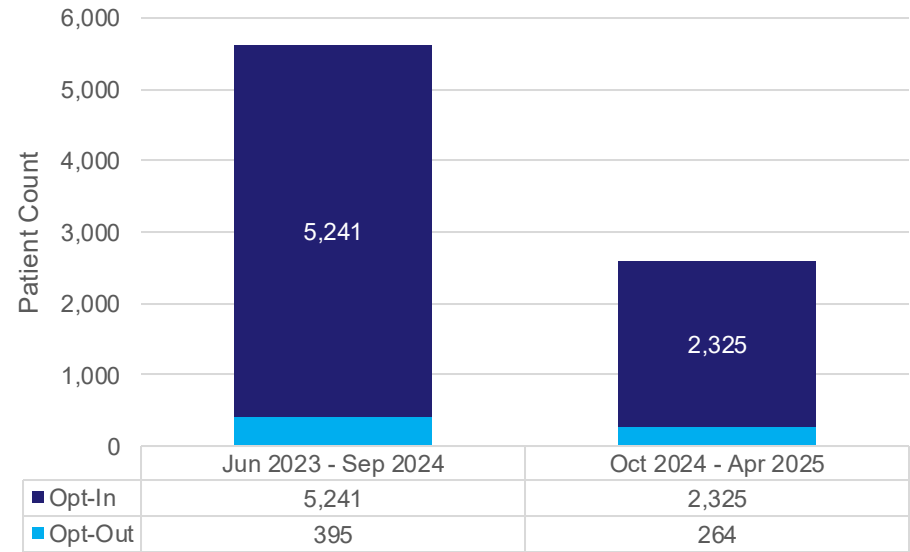
# Use of Epic for Clinical Trial Recruitment

**MyChart Recruitment and Clinical Trial Alerts are two Epic features for expanding clinical trial recruitment**

MyChart Recruitment	Clinical Trial Alerts
<ul style="list-style-type: none"><li>• Individuals are identified as potentially eligible for a clinical trial via data stored in the electronic health record</li><li>• Patient is sent a MyChart message to alert them of potential eligibility in a study</li><li>• The patient can express or decline interest in the study via MyChart</li></ul>	<ul style="list-style-type: none"><li>• Individuals are identified as potentially eligible for a clinical trial via data stored in the electronic health record</li><li>• The provider receives an alert that a specific patient may be eligible for a study</li><li>• Via Epic, the provider can alert the study team if the patient is interested in participating in a study</li></ul>

	MyChart Recruitment	Clinical Trial Alerts
Live	13	8
In Progress	5	1

# Metrics for MyChart Research Opt-Out



- In total 8,225 patients have responded to the Epic Research Consent, with 2,589 new responses since October 2024
- 659 (8.0%) of respondents have opted-out of being contacted via MyChart for research studies suggested by information in their electronic health record

# **MSDW Roadmap**

## **May 2025 –**

## **November 2025**

# MSDW Projects in Progress

	Project	Target Date	New Capabilities for Researchers
1	<b>Announce the ability to Identify patients with select somatic genetic mutations in Leaf</b>	2025-Q2	Enable researchers to obtain counts of patients with expanded list of somatic genetic mutations via Leaf
2	<b>Explore and prototype new approaches extract data Pathology Notes using AI</b>	2025-Q3	Facilitate medical research by providing quick access to relevant pathology data

# MSDW Resources & Useful Links

# MSDW Resources and Useful Links



Open a ticket: <https://labs.icahn.mssm.edu/msdw/open-a-ticket/>



Training and Tutorials: <https://labs.icahn.mssm.edu/msdw/training/>



About Leaf: <https://labs.icahn.mssm.edu/msdw/about-leaf/>



About ATLAS: <https://labs.icahn.mssm.edu/msdw/about-atlas/>



About TriNetX: <https://live.trinetx.com/>



Contact Us: <https://labs.icahn.mssm.edu/msdw/contact-us/>



Frequently Asked Questions: <https://labs.icahn.mssm.edu/msdw/faqs/>

# Thank You