



Leveraging AI to Advance Public Health Data Infrastructure

Aug. 18, 2025 | 2025 ARC Preconference

SHAPING TOMORROW'S PUBLIC HEALTH TODAY.

Session Objectives

- Understand the foundational concepts of AI and how it can be applied to public health.
- Explore real-world applications of GenAI in public health through case studies.
- Recognize challenges and best practices in implementing GenAI solutions in public health settings.

Agenda

1. Understanding AI in Public Health
2. Speaker Introductions
3. PHA Speaker presentations showcasing GenAI being used in Public Health
4. Q&A

Understanding AI in Public Health

Key Definitions

Artificial Intelligence (AI) is the development, implementation, and use of computer systems that can perform tasks that typically require human intelligence. AI describes technology that makes computers (seem to) act rationally.

Machine learning (ML) allows a computer to analyze data to do a task without being explicitly programmed. Common functions of machine learning are to (1) find patterns, like groupings of similar items and (2) to guess or predict an output based on a set of inputs.

Narrow AI also known as “weak” or “traditional” AI, focuses on performing specific tasks within a limited domain, such as image recognition, speech synthesis, or playing chess. Narrow AI has been in use for decades (decision support, google searches)

General AI refers to highly autonomous systems that feel like they possess human-level intelligence and can handle various cognitive tasks across different domains. Large language models (LLMs) are the major advancement in general AI.

- **Generative AI (GenAI)** refers to artificial intelligence systems that can create new, original content, such as text, images, or music, by learning patterns and structures from existing data.



Key Definitions

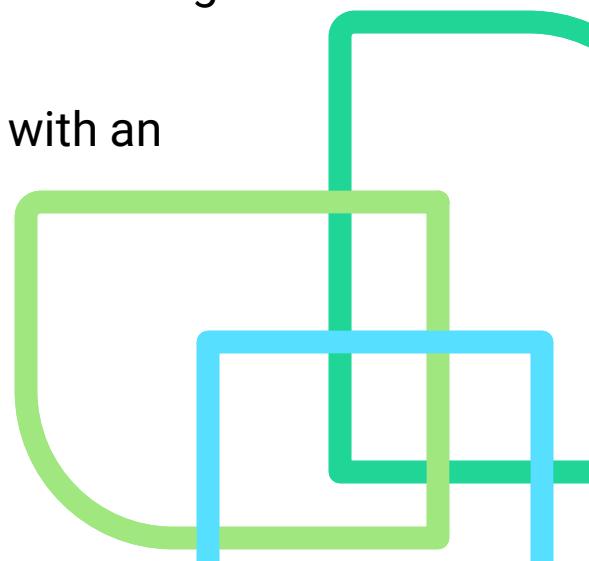
Large language models: These are advanced AI models that can understand and generate human-like text based on the patterns and structures they learn from vast amounts of data

Text chunking and vectorization/vector store: This involves breaking down text into smaller chunks and converting them into numerical vectors that can be stored and processed by AI models

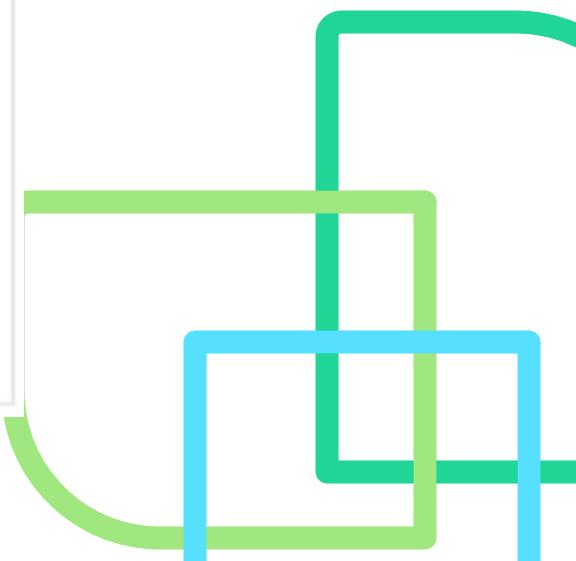
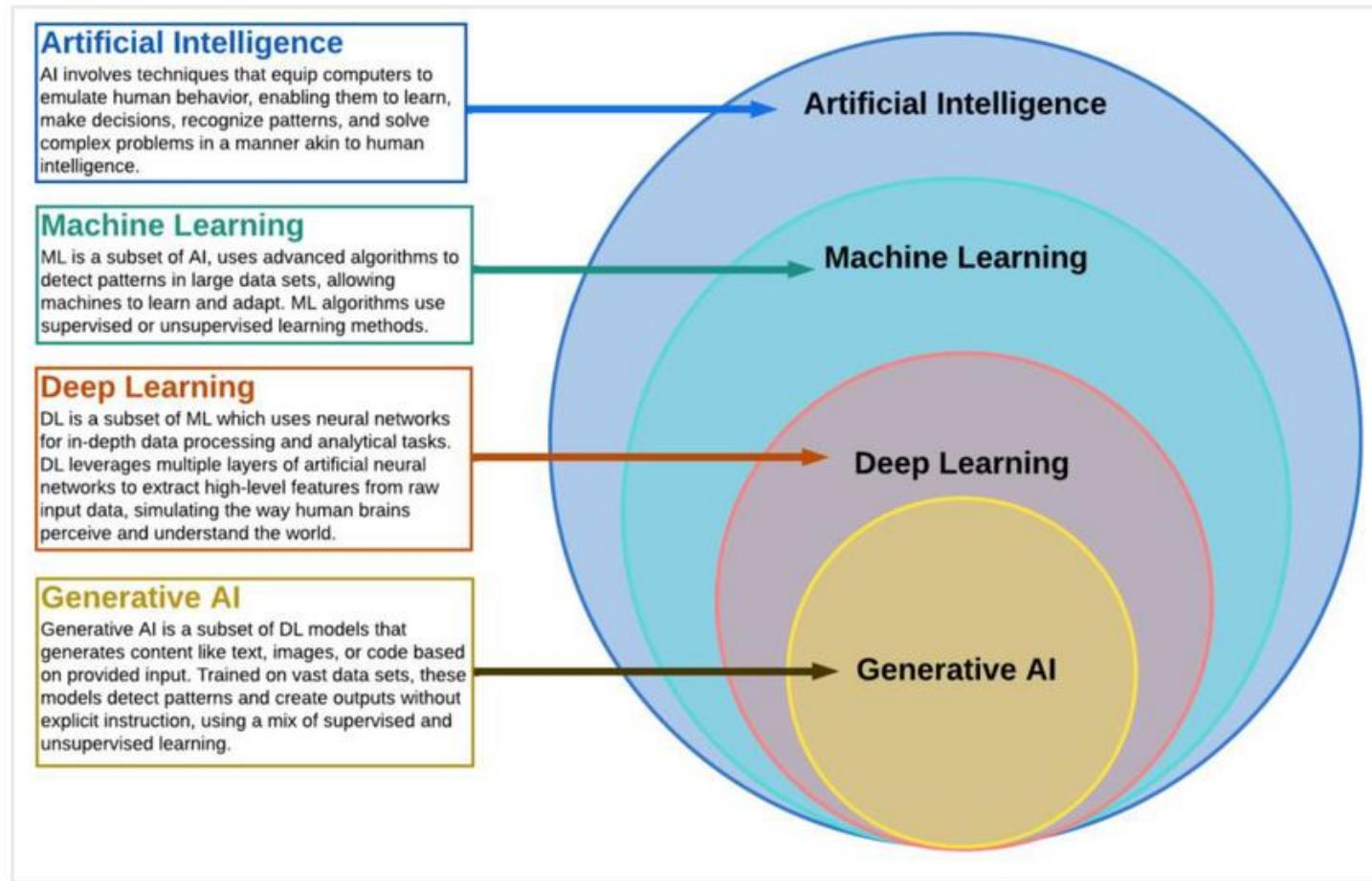
Retrieval augmented generation (or RAG) approach: This is a method that combines the strengths of retrieval-based and generation-based models to produce more accurate and contextually relevant response

Deterministic variables: These are variables in a model that have a fixed value and do not change based on probability

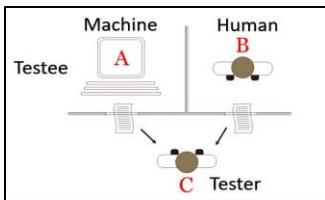
Probabilistic variables: These are variables that have a range of possible values, each with an associated probability



A Comparative View of AI, ML, Deep Learning, & GenAI



A Comparative View of AI, ML, Deep Learning, & GenAI



Turing Test (1950)

A vision of what's to come



IBM Watson (2011)

AI defeats humans in Jeopardy!



OpenAI's GPT-3 Release (2020)

AI is capable of generating human-like text



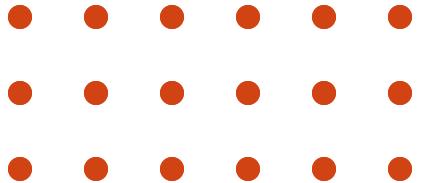
Proliferation of genAI (2023)

GenAI has expanded across industries and many people use AI daily



America's AI Action Plan (2025)

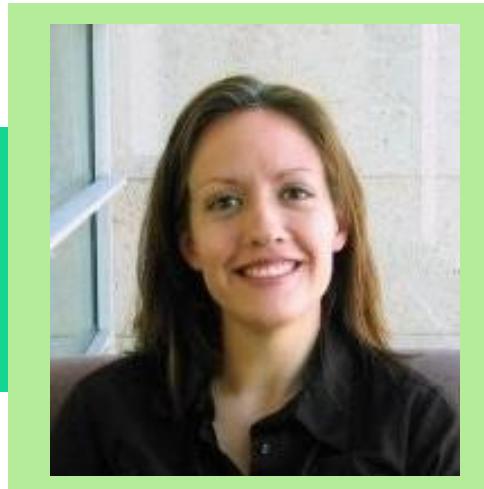
White House unveiled comprehensive AI Strategy in July of 2025



Meet the Speakers

**Dejan Jovanov**

Chief Data & Informatics Officer,
Division of Patient Safety & Quality,
Office of Policy Planning & Statistics
Illinois Department of Public Health

**Stephanie Meyer, MPH**

Epidemiology & Data Unit Supervisor,
Emerging Infectious Diseases
Epidemiology & Response Section,
Infectious Disease Epidemiology,
Prevention, & Control Division
Minnesota Department of Health

**Jim Daniel, MPH**

Leader of State and Local
Public Health at Amazon
Web Services (AWS)



Measles Outbreak Simulator Dashboard

Changing the response from reactive to proactive



Dejan Jovanov (he/him)
Chief Data and Informatics Officer
Illinois Department of Public
Health

Setting the Stage:

Purpose:

Projects and estimates the likelihood of measles outbreaks in Illinois schools (PreK–12) using:

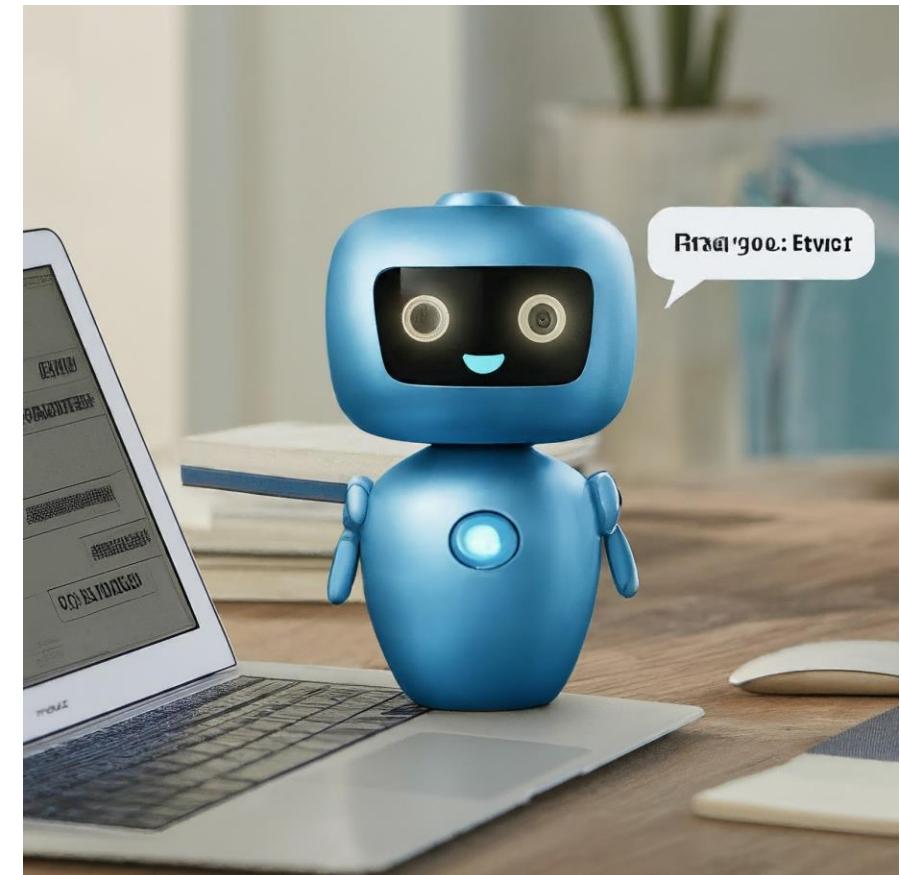
- School vaccination rates
- Enrollment data
- Additional model parameters

• Background:

- Released by the Illinois Department of Public Health (IDPH) in April 2025 due to a nationwide surge in measles cases
- Inspired by the University of Texas Austin *epiEngage Measles Outbreak Simulator*
- Part of our initiative to transform the IDPH data ecosystem from reactive to proactive

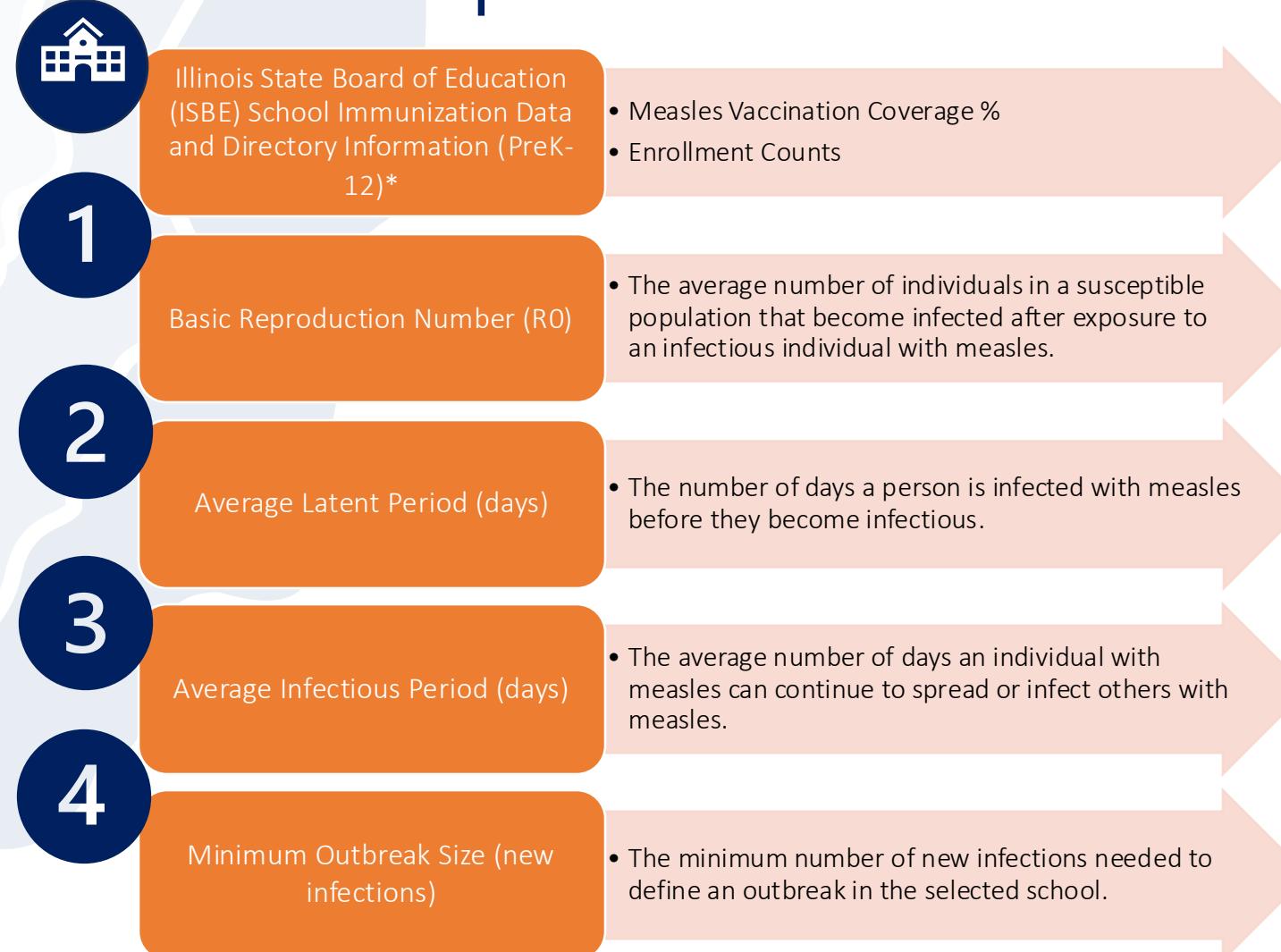
• Use:

- Supports school administrators and staff in outbreak prevention and control efforts
- Inform the public



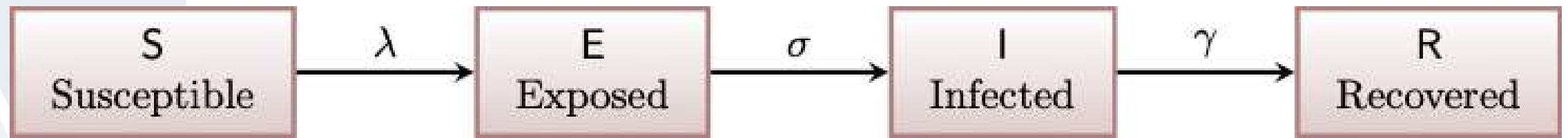
Measles Simulator Dashboard

Data Components



Likelihood and Size of a Measles Outbreak in an Illinois School

SEIR model used for Measles Prediction



- **Model Overview:**
- **SEIR** = *Susceptible* \rightarrow *Exposed* \rightarrow *Infected* \rightarrow *Removed*
- Population moves between compartments based on defined rates.
- **Key Parameters:**
- **λ (Exposure Rate):** How quickly susceptible individuals become exposed
- **σ (Latent Rate):** How quickly exposed individuals become infected
- **γ (Removed Rate):** How quickly infected individuals recover or are removed
- Also includes: vaccine rate, initial infection, contact rate, basic reproduction number (R_0)
- Defaults follow CDC Measles Clinical Diagnosis Fact Sheet

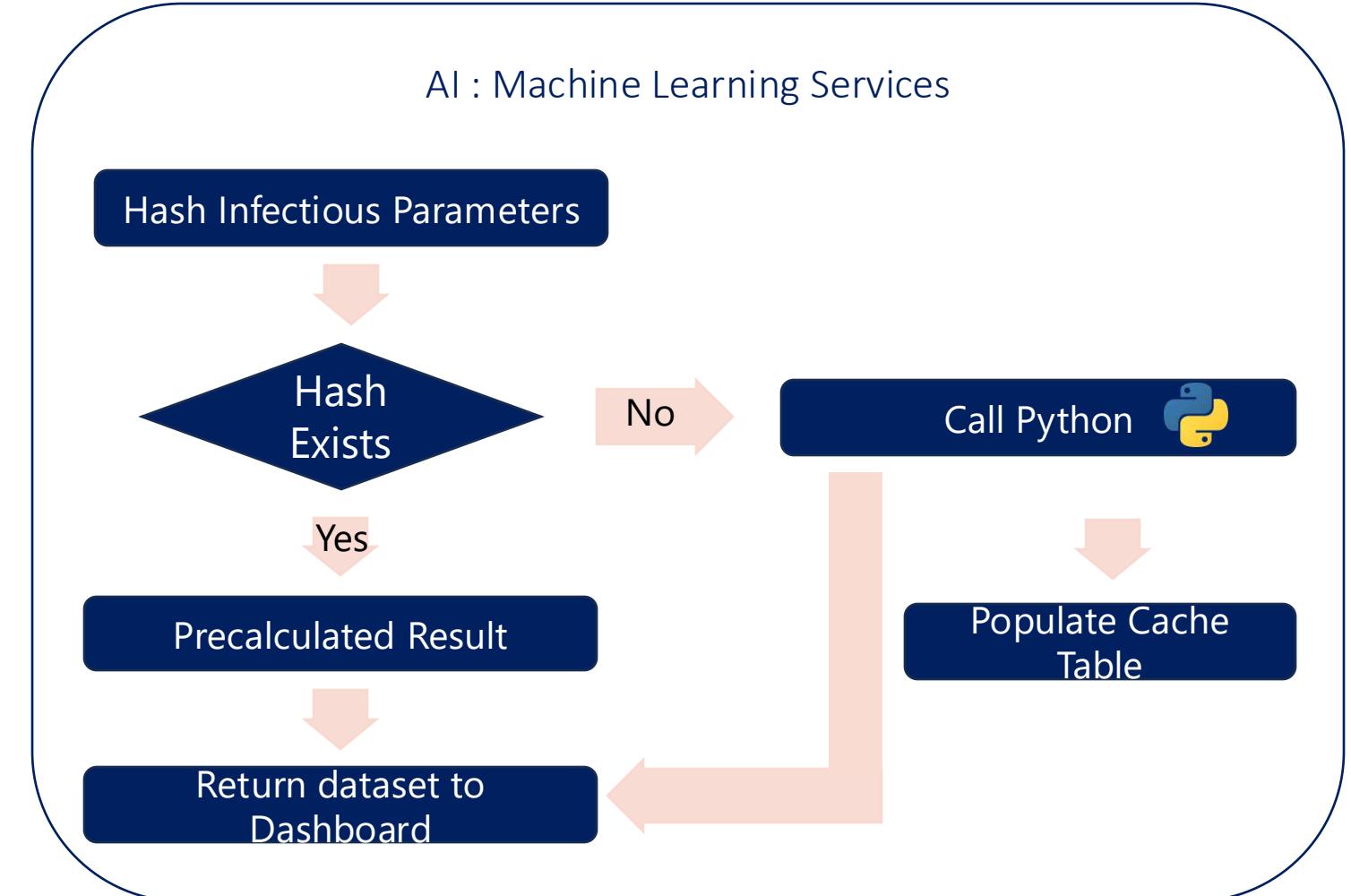
Measles Simulator Dashboard Calculation Process



Request
from
Dashboard

Infectious Parameters

- Precalculated schools with under 95% vaccination rate with default parameters
- Total of 7,374 cached results
- Peak of 2,488 new cached results on 4/9/25 when the dashboard was on the news
- Other options:
 - TabPy
 - Snowflake



Measles Simulator Dashboard

Dashboard Parameters



1. Choose Enrollment & Vaccination Rates

- Select manually or pick a specific school.

1

Measles Outbreak Projections in Illinois Schools

Select the county, type of school funding and school district you want to see data for.
If no results are returned, confirm that the school district filter is either set to (All), or to a relevant school district in the selected county-school funding combination.

County	Funding Type	School District
<input type="text" value="City of Chicago"/>	<input type="text" value="Public"/>	<input type="text" value="(All)"/>

Select a school from the list of relevant schools below to generate a simulation.

School District	Facility Name	Enrollment PreK-12	School Vaccination Rate
Burnham SD 154-5	Burnham Elem School	146	98.6%
Calumet Public SD 132	Burr Oak Academy	318	95.3%
	Burr Oak Elem School	288	97.2%
	Calumet Elem School	309	98.1%

2. Filter Schools

- Use County, Funding Type, and School District filters.
- If the county is changed, adjust the School District drop-down or set to (All).

2

3. Update & Select

- The table updates based on your selections.
- Click a school to use its enrollment and vaccination rate.
- Values can also be adjusted manually to model changes.

3

School Enrollment	School Vaccination Rate (%)	Students Initially Infected
<input type="text" value="500"/>	<input type="text" value="85.0%"/>	<input type="text" value="1"/>

You can change the default epidemic parameters using the sliders below to generate a more customized projection.

Please note that the four parameters below are for use of local health departments and epidemiologists.
Interpretation of the information based on any adjustments to the epidemic parameters below is extremely nuanced and subject to misinterpretation.

Basic Reproduction Number (R0)	Average Latent Period (days)	Average Infectious Period (days)	Minimum Outbreak Size (new infections)
<input type="text" value="15"/>	<input type="text" value="10.5"/>	<input type="text" value="5"/>	<input type="text" value="10"/>

4. Adjust Epidemic Parameters

- Modify the minimum outbreak size to change the threshold for potential outbreaks.
- Other epidemic parameters can be updated for custom scenarios.

4

Chance of exceeding 10 new infections

58%

with selected model inputs and epidemic parameters

Likely outbreak size

45 - 72 cases

if there are over 10 new infections

Measles Simulator Dashboard

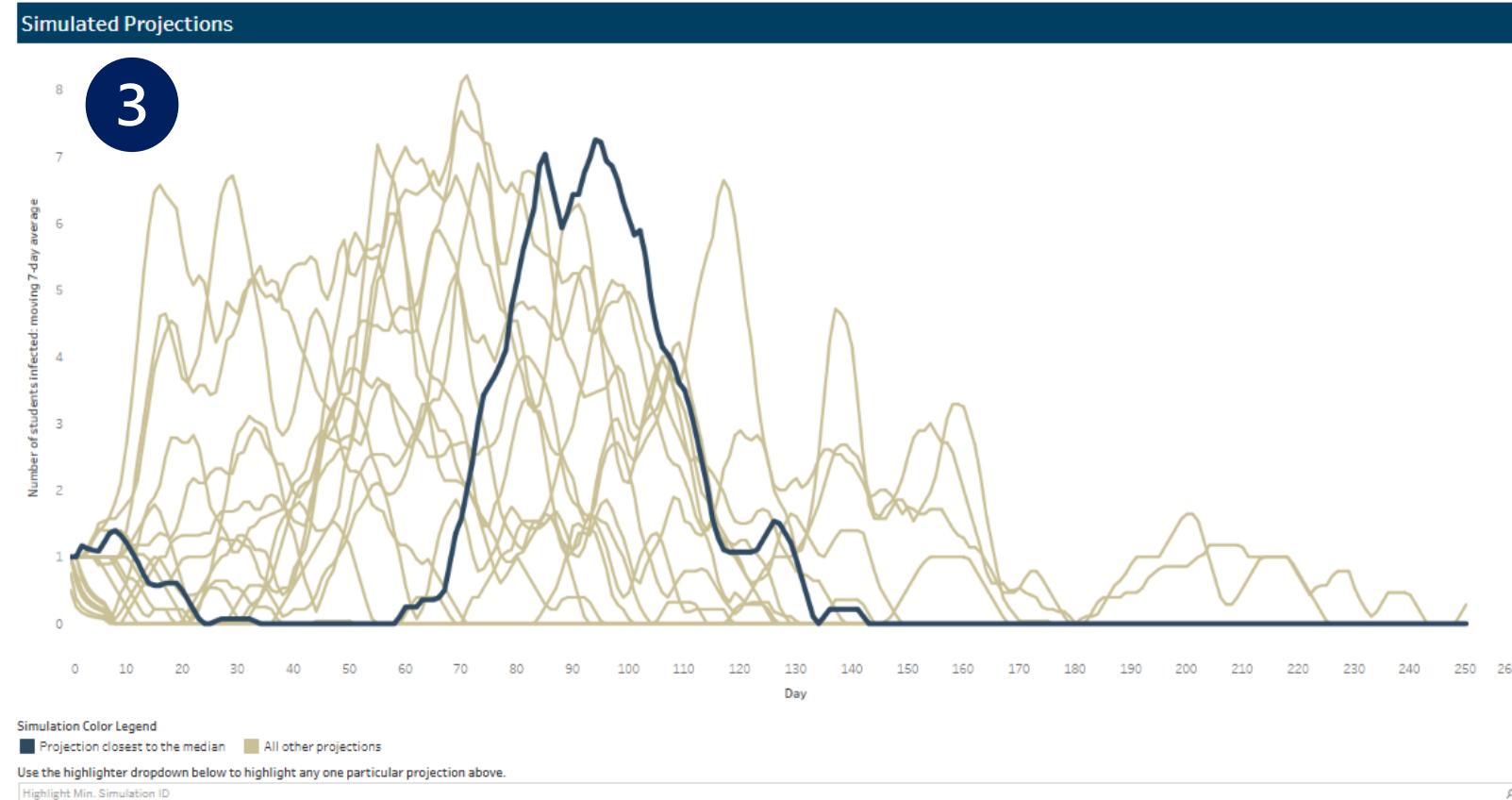
Dashboard Results

The results are calculated by running 200 stochastic simulations using the entered parameters.

1. The "Chance of exceeding X new infections" is the percentage of simulations that had more infections than the selected minimum (10 by default).
2. Likely outbreak size uses the middle 95% of the outbreaks that are larger than the selected minimum (10 by default).
3. The simulated projections show a random selection of 20 outbreak projections. The highlighted line is the projection with the outbreak size nearest to the median.

1	Chance of exceeding 10 new infections 58% with selected model inputs and epidemic parameters	2	Likely outbreak size 45 - 72 cases if there are over 10 new infections
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Projections and estimates for outbreaks provided below, are specific to the school population selected and incorporate the selected school vaccination coverage, enrollment, and additional model parameter values.
The calculation of the proportion of the school population susceptible to measles does not account for community immunity outside schools, interventions, or breakthrough infections amongst vaccinated students.



What's Next



- Adapt the SEIR model (adding compartments, age groups, vaccination status, seasonality) to fit the specific disease.
- Expand the use of the model on diseases with a clear incubation period (e.g., measles, COVID-19, influenza).
- Build additional dashboards



Questions?

Contact:

Dejan Jovanov: Dejan.Jovanov@Illinois.gov



Project CLAIRE: Comprehensive Language Model Artificial Intelligence for Record Evaluation - Assessing the use of Artificial Intelligence for Infectious Disease Surveillance

Stephanie Meyer, MPH

Epidemiology Supervisor | Infectious Disease Epidemiology, Prevention, and Control Division
Minnesota Department of Health

Background

- COVID-19 is one of many reportable diseases in Minnesota
- Minnesota is part of the CDC Emerging Infections Program
- Medical chart abstractions are often necessary
- COVID-NET monitors laboratory-confirmed, COVID-19-associated hospitalizations among children and adults

Reportable Diseases, MN Rule 6605.7000 & 6605.7000
651-201-5414 or 1-877-676-5414 24 hours a day, 7 days a week

Diseases Reportable to the Minnesota Department of Health

REPORT IMMEDIATELY BY TELEPHONE

- Anthrax (Bacillus anthracis) (●)
- Botulism (Clostridium botulinum) (●)
- Brucellosis (Brucella spp.) (●)
- Cholera (Vibrio cholerae) (●)
- Diphtheria (Corynebacterium diphtheriae) (●)
- Flea-borne typhus infection (●)
- (including at least: *Acaranthomoeba* spp., *Nieogliobia fowleri*, *Balamuthia* spp., *Sappinia* spp.) (●)
- Granuloma (Burkholderia mallei) (●)
- Hantavirus (Leptospiral spp.) (●)
- Measles (rubella) (●)
- Melioidosis (Burkholderia pseudomallei) (●)
- Meningococcal disease (Neisseria meningitidis) (invasive) (●)
- Middle East Respiratory Syndrome (MERS) (●)
- Orthopox virus (●)
- Plague (Yersinia pestis) (●)
- Poliomyelitis (●)
- Q fever (Coxiella burnetii) (●)
- Rabies (Lyssavirus) (●)
- Rubella and congenital rubella syndrome (●)
- Severe Acute Respiratory Syndrome (SARS) (●)
- Smallpox (variola) (●)
- Tularemia (Francisella tularensis) (●)
- Unusual or increased case incidence of any suspect infectious illness (●)
- Viral hemorrhagic fever (●)
- (including but not limited to Ebola virus disease and Lassa fever)

REPORT WITHIN ONE WORKING DAY

- Amebiasis (*Entamoeba histolytica*/diaper) (●)
- Anaplasmosis (*Anaplasma phagocytophili*) (●)
- Arbovirus (●)
- (including, but not limited to: E. Crosse encephalitis, eastern equine encephalitis, western equine encephalitis, St. Louis encephalitis, West Nile virus disease, Powassan virus disease, and Jamestown Canyon virus disease) (●)
- Babesiosis (*Babesia microti* spp.) (●)
- Bacillary angiomatosis (*Ureaplasma urealyticum*) (●)
- Campylobacteriosis (*Campylobacter* spp.) (●)
- Conjunctivitis (●)
- Carbapenem-resistant Enterobacteriaceae (CRE) (●)
- Cat scratch disease (infection caused by *Bartonella* species) (●)
- Chancroid (Neisseria ulcerans) (●)
- Chikungunya virus disease (●)
- Chlamydia trachomatis infections (●)
- Coccidioidomycosis (●)
- Granulocystis (Granulocystis kutscheri) in infants under one year of age (●)
- Cryptosporidiosis (*Cryptosporidium* spp.) (●)
- Cyclosporiasis (*Cyclospora* spp.) (●)
- Davidson's disease (●)
- Diphyllobothriosis (liver) infection (●)
- Escherichia (Escherichia spp.) (●)
- Esopatidis (caused by viral agent) (●)
- Enteric Escherichia coli infection (●)
- (*E. coli* O157:H7, other Shiga-toxin-producing *E. coli*, enteropathogenic *E. coli*, enterotoxigenic *E. coli*, enteroinvasive *E. coli*, enterohemorrhagic *E. coli*, enteropathogenic *E. coli*, or other pathogenic *E. coli*) (●)
- Giardiasis (*Giardia intestinalis*) (●)
- Gonorrhoea (*Neisseria gonorrhoeae* infections) (●)
- Hemophagocytic lymphohistiocytosis (HLH) (●)
- Hantavirus infection (●)
- Hepatitis A, B, C, D, E, G, and J virus (including A, B, C, D, and E) (●)
- Histoplasmosis (*Histoplasma capsulatum*) (●)
- Human immunodeficiency virus (HIV) infection, including Acquired Immunodeficiency Syndrome (AIDS) (●)
- Influenza (●)
- (including case incidence, critical illness, or laboratory-confirmed cases) (●)
- Kawasaki disease (●)
- Kingella spp. (invasive only) (●)
- Legionellosis (*Legionella* spp.) (●)
- Leprosy (*Hansen's disease*) (*Mycobacterium leprae*) (●)
- Leptospirosis (*Leptospira interrogans*) (●)
- Listeriosis (*Listeria monocytogenes*) (●)
- Lyme disease (*Borrelia burgdorferi* and other *Borrelia* spp.) (●)
- Malaria (*Anopheles* spp.) (●)
- Meningitis (caused by viral agent) (●)
- Murine (●)
- Neonatal sepsis (●)
- (bacteria isolated from a sterile site, excluding coagulase-negative Staphylococcus) (●)
- Neonatal sepsis (caused less than seven days after birth) (●)
- Perimeningitis (*Escherichia coli*) (●)
- Pneumocystis (*Pneumocystis jirovecii*) (●)
- Retovirus infections (●)
- Salmonellosis, including typhoid (*Salmonella* spp.) (●)
- Shigellosis (*Shigella* spp.) (●)
- Spotted fever group rickettsiosis (●)
- (Rocky Mountain spotted fever, including Rocky Mountain spotted fever) (●)
- Staphylococcus aureus (●)
- (only vancomycin-intermediate Staphylococcus aureus (VISA), vancomycin-resistant Staphylococcus aureus (VRSA), or critical illness due to community-associated Staphylococcus aureus in a previously healthy individual) (●)
- Streptococcal disease - invasive disease caused by Groups A and B streptococci and *S. pneumoniae* (●)
- Streptococcal disease - non-invasive *S. pneumoniae* (●)
- Streptococcal pneumonia (●)
- Syphilis (*Treponema pallidum*) (●)
- Tetanus (*Clostridium tetani*) (●)
- Toxic shock syndrome (●)
- Toxoplasmosis (*Toxoplasma gondii*) (●)
- Transmissible spongiform encephalopathy (●)
- Tuberculosis (*Mycobacterium tuberculosis* complex) (●)
- (pulmonary or extrapulmonary sites of disease, including clinically diagnosed disease). Latent tuberculosis infection is not reportable.
- Typhus (*Ixodes ticks*) (●)
- Unusual deaths and unexplained critical illness (possibly due to infectious cause) (●)
- Varicella (*Varicella*) (●)
- Vibrio spp. (●)
- Yellow fever (●)
- Yersiniosis (enteric *Yersinia* spp., regardless of specimen source) (●)
- Zoster (shingles) (●)
- (all cases 18 years old; unusual case incidence/complications regardless of age)

SENTINEL SURVEILLANCE

Diseases reportable through sentinel surveillance are reportable based on the residence of the patient or the healthcare facility. Sentinel surveillance is not statewide reporting.

- Staphylococcus aureus (●)
- Candidemia (*Candida* spp.) (●)
- Carapenem-resistant Acinetobacter spp. (CRAB), including carbapenemase-producing (CR-PA) (●)
- Clostridium difficile (●)
- Respiratory syncytial virus (RSV) (●)
- Non-tuberculous Mycobacteria (NTM), pulmonary and extrapulmonary (●)

FOOTNOTES

- Submission of clinical materials required. Submit isolates or, if an isolate is not available, submit material containing the infectious agent in the following order of preference: blood, tissue, sputum, urine, or other non-sterile material. Call the MDH Public Health Laboratory at 651-205-4951 for instructions.
- Invasive disease only. Isolated from a normally sterile site, e.g.: blood, CSF, joint fluid, etc.
- Report cases of SARS in another agency's respiratory network, also report cases of health care workers hospitalized for pneumonia or acute respiratory distress syndrome.
- Also report a pregnancy in a person with ZIKV, or a person chronically infected with hepatitis B, HCV, or syphilis.

TO REPORT

- For immediate reporting call: 651-201-5414 or 1-877-676-5414.
- Report forms can be downloaded at www.health.state.mn.us/ideas/report.

44-57153 | 10/2019

What is a Medical Chart Review?

ED Vitals:

Patient Vitals for the past 24 hrs:

	BP	Temp	Pulse	Resp	SpO2	Height	Weight
09/02/23 0202	127/85	99 °F (37.2 °C)	97	22	94 %	—	—
09/02/23 0000	(!) 115/91	—	(!) 110	(!) 34	95 %	—	—
09/01/23 2345	128/78	—	(!) 103	(!) 32	95 %	—	—
09/01/23 2330	—	—	(!) 103	(!) 26	95 %	—	—
09/01/23 2315	(!) 128/116	—	(!) 108	(!) 32	94 %	—	—
09/01/23 2306	136/88	99.5 °F (37.5 °C)	(!) 107	20	97 %	5' 4.02" (1.626 m)	84.8 kg

MDM:

██████████ is a 76 y.o. female who presents for the above. The differential includes pneumonia, sepsis, viral syndrome, antibiotic reaction. The patient has generalized weakness here and recent diagnosis of pneumonia. Patient is given albuterol treatment. Sepsis protocol was started and she is given some lactated Ringer's. Patient is given levofloxacin after extensive conversation with her about her allergies and given the pneumonia. Chest x-ray does show significant left-sided infiltrate. She has significant hyponatremia as well. I did discuss the case with renal on-call who recommended fluid resuscitation and recheck of sodium. Patient is transferred to hospitalist service for further work-up and care.

DIAGNOSIS:

ICD-10-
CM

1. Pneumonia of left lung due to infectious organism, unspecified part of lung J18.9
2. Hyponatremia E87.1

Medical charts are read and reviewed to determine:

- Course of illness
- Course of hospitalization
- Treatments
- Underlying Conditions

I Notes (group 1 of 2) (continued)

g: Take 25 mg by mouth once daily.					
colecalciferol, vitamin D3, 25 mcg, 1000 unit, 25 mcg (1,000 unit) oral tablet	9/29/2022	Patient	Yes	Yes	
g: Take 25 mcg by mouth once daily.					
clobenzaprine (FLEXERIL) 10 mg oral tablet	9/29/2022	Patient	Yes	Yes	
g: Take 10 mg by mouth every evening.					
ulaglutide (TRULICITY) 1.5 mg/0.5 mL SubQ	9/23/2022	Patient	Yes	Yes	
g: Inject 1.5 mg under the skin every 7 (seven) days.					
abapentin (NEURONTIN) 300 mg oral capsule	9/29/2022	Patient	Yes	Yes	
g: Take 300 mg by mouth three times a day.					
ipiZIDE (GLUCOTROL XL) 5 mg oral extended release tablet 24 HR	9/29/2022	Patient	Yes	Yes	
g: Take 5 mg by mouth twice a day.					
multivit-mins/iron/folic/lycop (CENTRUM EN ORAL)	9/29/2022	Patient	Yes	Yes	
g: Take 1 capsule by mouth once daily.					
aloxone (NARCAN) 4 mg/actuation Nasal spray	PRN	Patient	Yes	Yes	
g: Instill 4 mg into one nostril as needed. Seek emergency care immediately after use.					
oxyCODONE (OXYCONTIN) 20 mg oral extended release tablet 12 HR	9/29/2022	Patient	Yes	Yes	
g: Take 20 mg by mouth twice a day.					
oxyCODONE, immediate release, (ROXICODONE) 15 mg oral tablet	9/29/2022	Patient	Yes	Yes	
g: Take 15 mg by mouth every 4 (four) hours as needed.					
antoprazole (PROTONIX) 40 mg oral delayed release tablet	9/29/2022	Patient	Yes	Yes	
g: Take 40 mg by mouth once daily.					
waroxaban (XARELTO) 20 mg oral tablet	9/30/2022 @0200	Patient	Yes	Yes	
g: Take 20 mg by mouth once daily.					
msulosin (FLOMAX) 0.4 mg oral capsule	9/29/2022	Patient	Yes	Yes	
g: Take 0.4 mg by mouth at bedtime.					
aZODone (DESYREL) 100 mg oral tablet	9/29/2022	Patient	Yes	Yes	
g: Take 100 mg by mouth at bedtime.					

Medical Chart Layout

- Medications or treatments may be listed over multiple pages or multiple days
- Each treatment may have been administered already or something that the patient will take home
- Details and nuance about dosage and timing can be very important in relation to an infection

Chart Abstractions

4. Height:	5. Weight:	6. BMI (non-pregnant cases and cases 2+ years old):
65 <input type="radio"/> in <input checked="" type="radio"/> cm reset	116.3 <input type="radio"/> lbs <input checked="" type="radio"/> kgs reset	42.7 View equation <input type="text" value="42.7"/>
7. Smoker (tobacco):		7a. Smoker (marijuana/vaping) [MN question]
<input checked="" type="radio"/> Current <input type="radio"/> Former <input type="radio"/> No/ Unknown reset		<input type="radio"/> Current <input type="radio"/> Former <input checked="" type="radio"/> No/ Unknown reset
Type of non-tobacco smoking (select all that apply):		
8. Environmental tobacco smoke exposure (for pediatric patients ≤12years):		
9. Alcohol misuse:		
<input type="radio"/> Current <input checked="" type="radio"/> Former <input type="radio"/> No/ Unknown reset		
10. Substance misuse:		
<input checked="" type="radio"/> Current <input type="radio"/> Former <input type="radio"/> No/Unknown reset		
11. Substance MisuseType (current use only) (Select all that apply):		
<input type="checkbox"/> IVDU <input checked="" type="checkbox"/> Opioids <input type="checkbox"/> Cocaine <input type="checkbox"/> Methamphetamine <input type="checkbox"/> Marijuana <input type="checkbox"/> Polysubstance abuse - not otherwise specified <input type="checkbox"/> Unknown <input type="checkbox"/> Other, specify		
12. Code status on admission:		
<input checked="" type="radio"/> Full code <input type="radio"/> DNR (Do Not Resuscitate) / DNI (Do not Intubate) / CMO (Comfort measures only) <input type="radio"/> Unknown reset		

ICU and Other interventions	
Invasive mechanical ventilation?	BiPAP or CPAP?
<input checked="" type="radio"/> Yes <input type="radio"/> No <input type="radio"/> Unknown reset	<input type="radio"/> Yes <input checked="" type="radio"/> No <input type="radio"/> Unknown reset
High flow nasal cannula (e.g. Vapotherm)?	Supplemental Oxygen?
<input type="radio"/> Yes <input checked="" type="radio"/> No <input type="radio"/> Unknown reset	
4. ECMO?	
<input type="radio"/> Yes <input checked="" type="radio"/> No <input type="radio"/> Unknown reset	
6. Renal Replacement Therapy (RRT) or Dialysis?	
Includes Peritoneal Dialysis (PD), Hemodialysis (HD), Continuous Venovenous Hemofiltration (CVVH), Continuous Venovenous Hemodialysis (CVVHD), and Slow Continuous Ultrafiltration (SCUF)	
<input type="radio"/> Yes <input checked="" type="radio"/> No <input type="radio"/> Unknown reset	
7. Was the patient admitted to an intensive care unit (ICU)?	
<input checked="" type="radio"/> Yes <input type="radio"/> No <input type="radio"/> Unknown reset	
7a. Date of ICU admission:	7b. Date of ICU discharge:
12-24-2024 M-D-Y	12-25-2024 M-D-Y
F. Outcome	
1. What was the outcome of the patient upon discharge?	
<input type="radio"/> Alive <input checked="" type="radio"/> Died during hospitalization <input type="radio"/> Unknown reset	
2. Date of death:	
12-25-2024 M-D-Y	

Medical Chart Review

- Manual chart review involves multiple staff
- COVID-NET includes a sample of hospitalized cases for chart review

Season*	2020-2021	2021-2022	2022-2023	2023-2024
Total COVID-NET Cases	39,318	34,154	14,908	12,971
Chart Reviews Completed	20,019 (50%)	7,006 (20%)	2,368 (16%)	2,669 (21%)
Review Staff	17	12	11	10

* Season starts October 1 through Sept 30 with exception of 2020-2021, which began in March 2020

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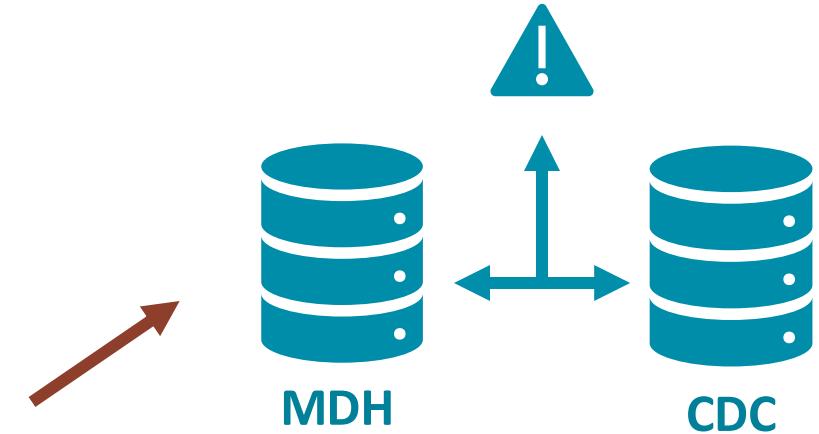
- The cost of full-time staff to review thousands of charts is not tenable

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Current State – Manual Process



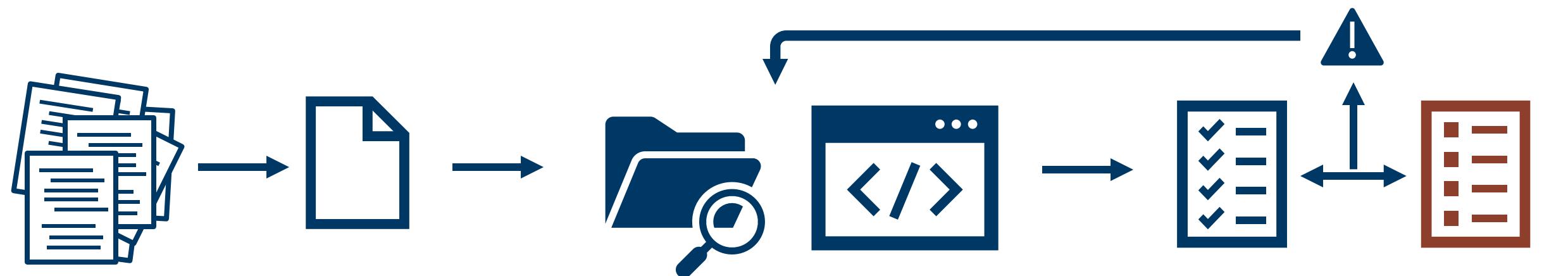
Manual medical record abstraction and data entry



Data sent from MDH to CDC, part of cooperative federal funding agreements

- If data entry errors are identified, problems are manually reviewed and re-abstracted

Future State



A document is generated with all the text from the medical record

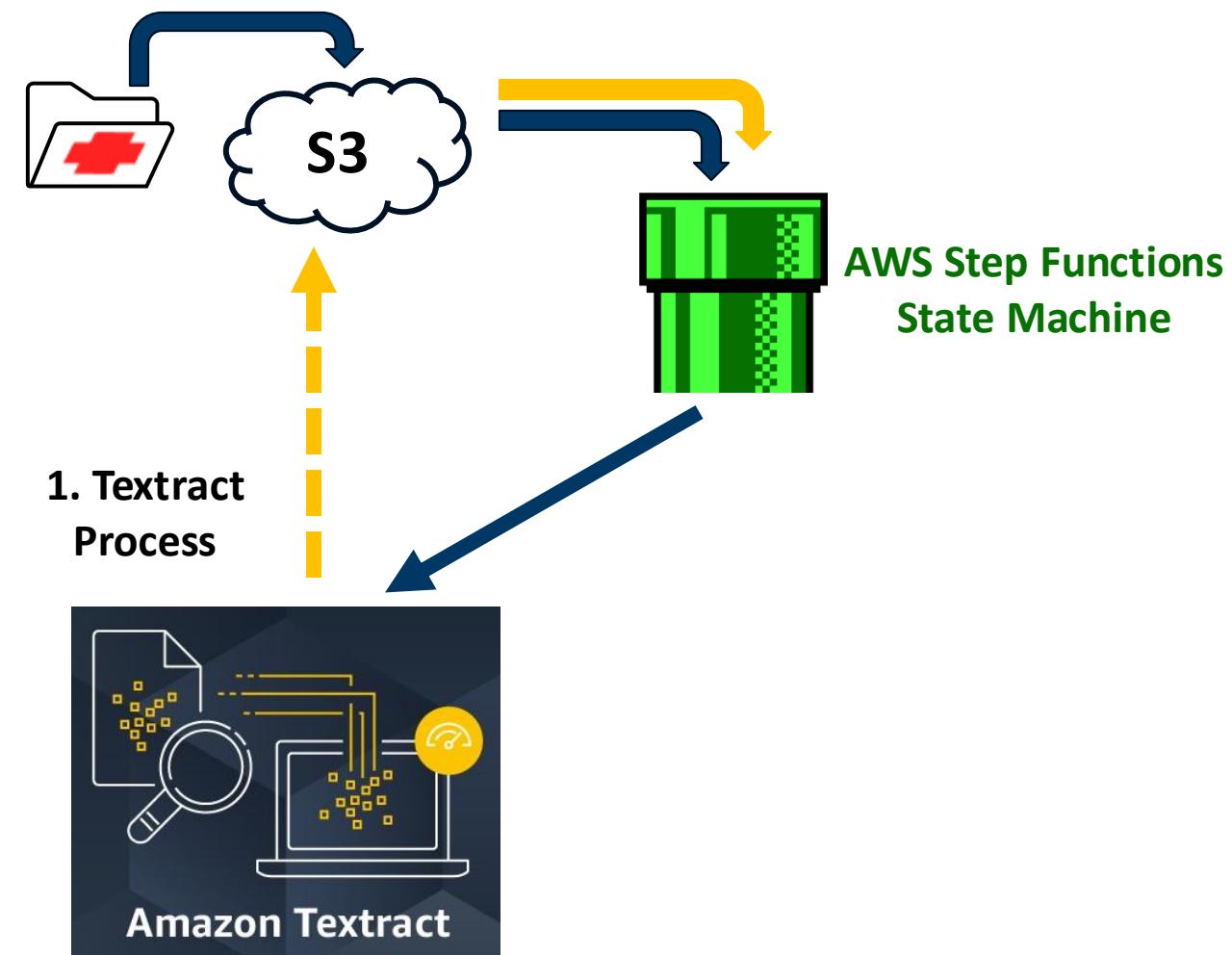
Use large language models to “read” the chart and extract key data elements
Includes code, prompt engineering, and large language model selection

Data is validated by comparison to manually collected data
Results are used to improve the abstraction pipeline

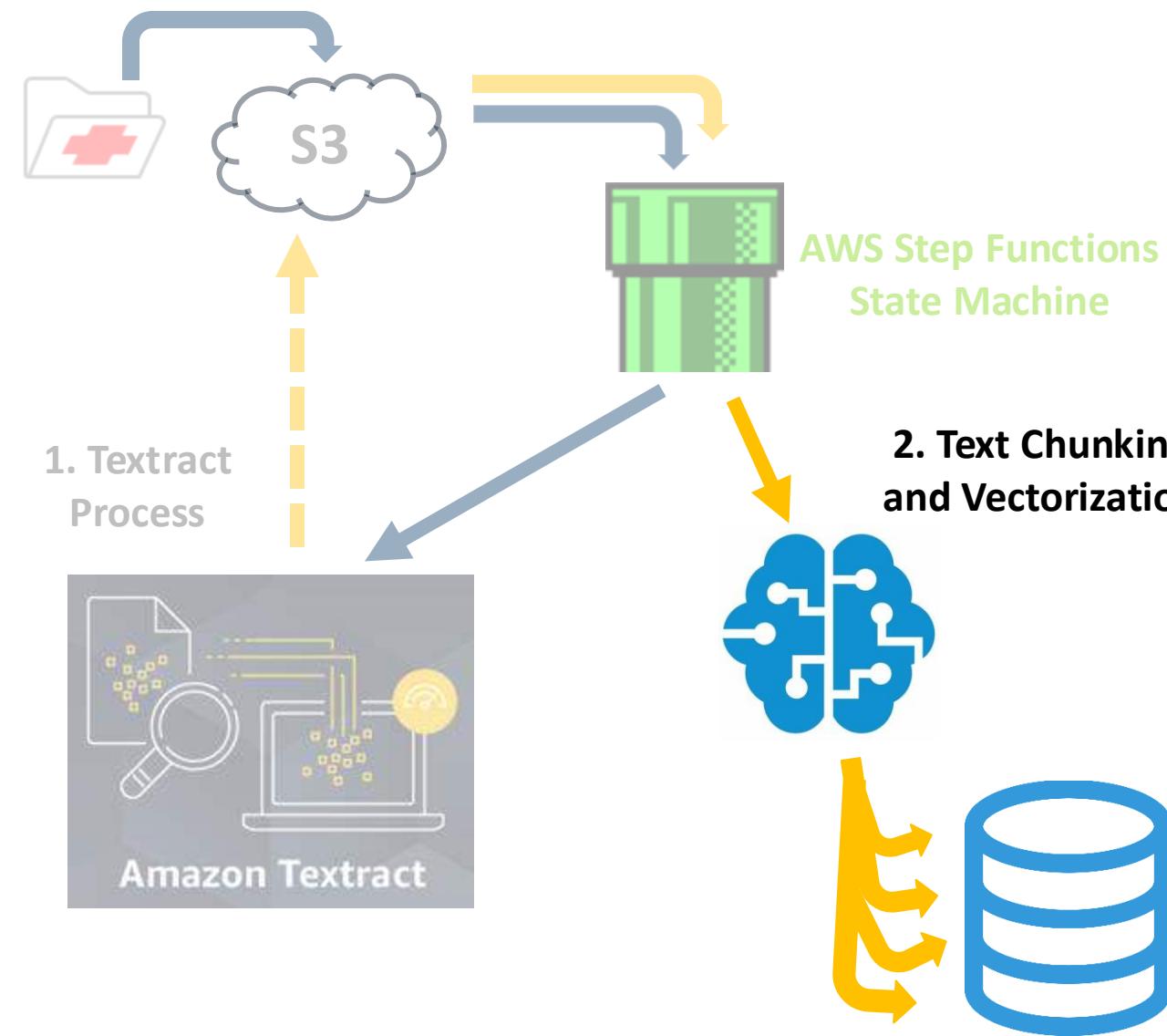
CLAIRE Pilot Project

- We identified 150 COVID-19 medical charts from one health system from the 2022-2023 respiratory season
- All charts were redacted PDF files
- All charts had been manually reviewed by Minnesota Department of Health (MDH) staff with answers entered into a REDCap database
- MDH partnered with MNIT MDH and Amazon Web Services (AWS) for evaluation of options
- Analysis of the results from text extracts of abstracted charts from the AWS pipeline were compared with manual abstractions

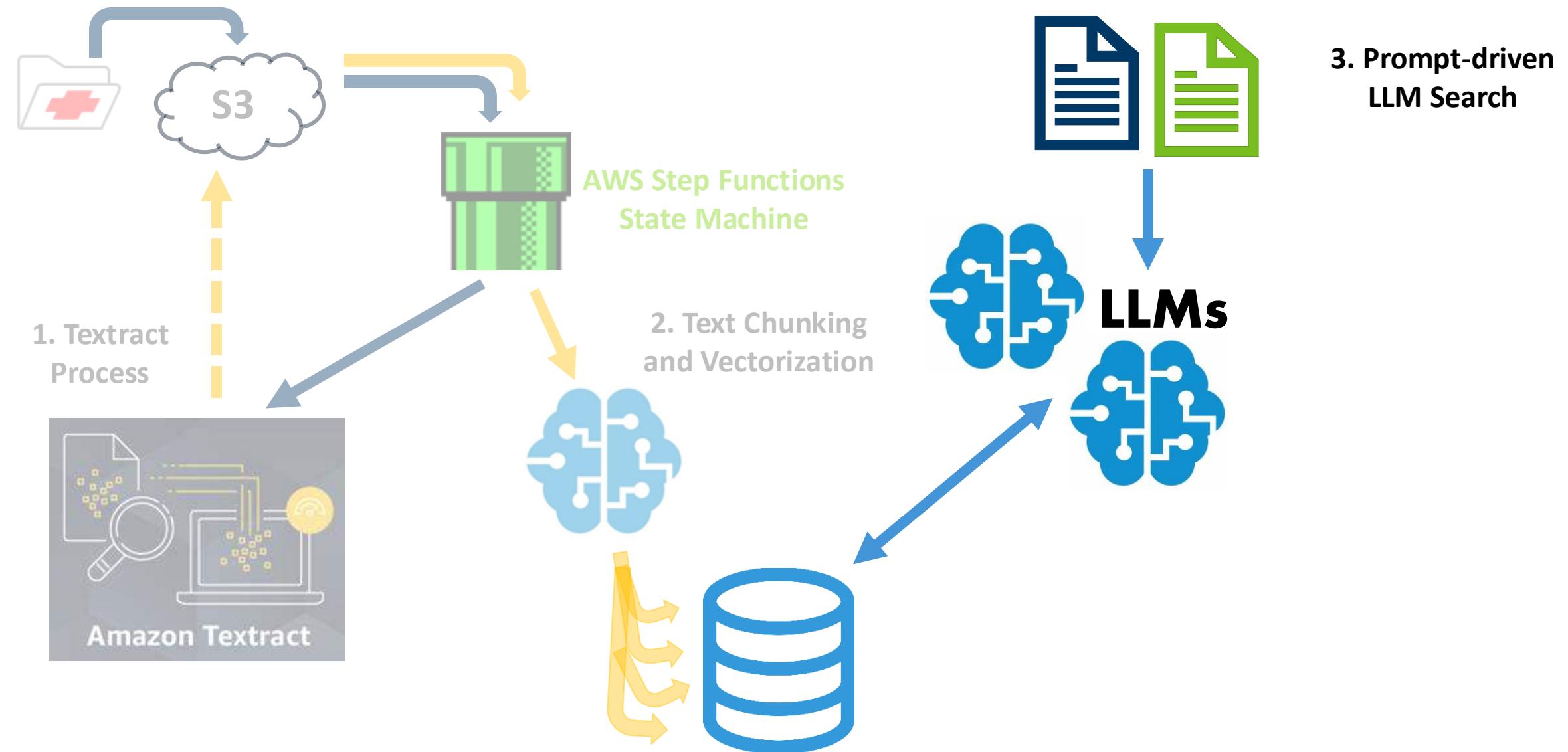
CLAIRE Pilot Project



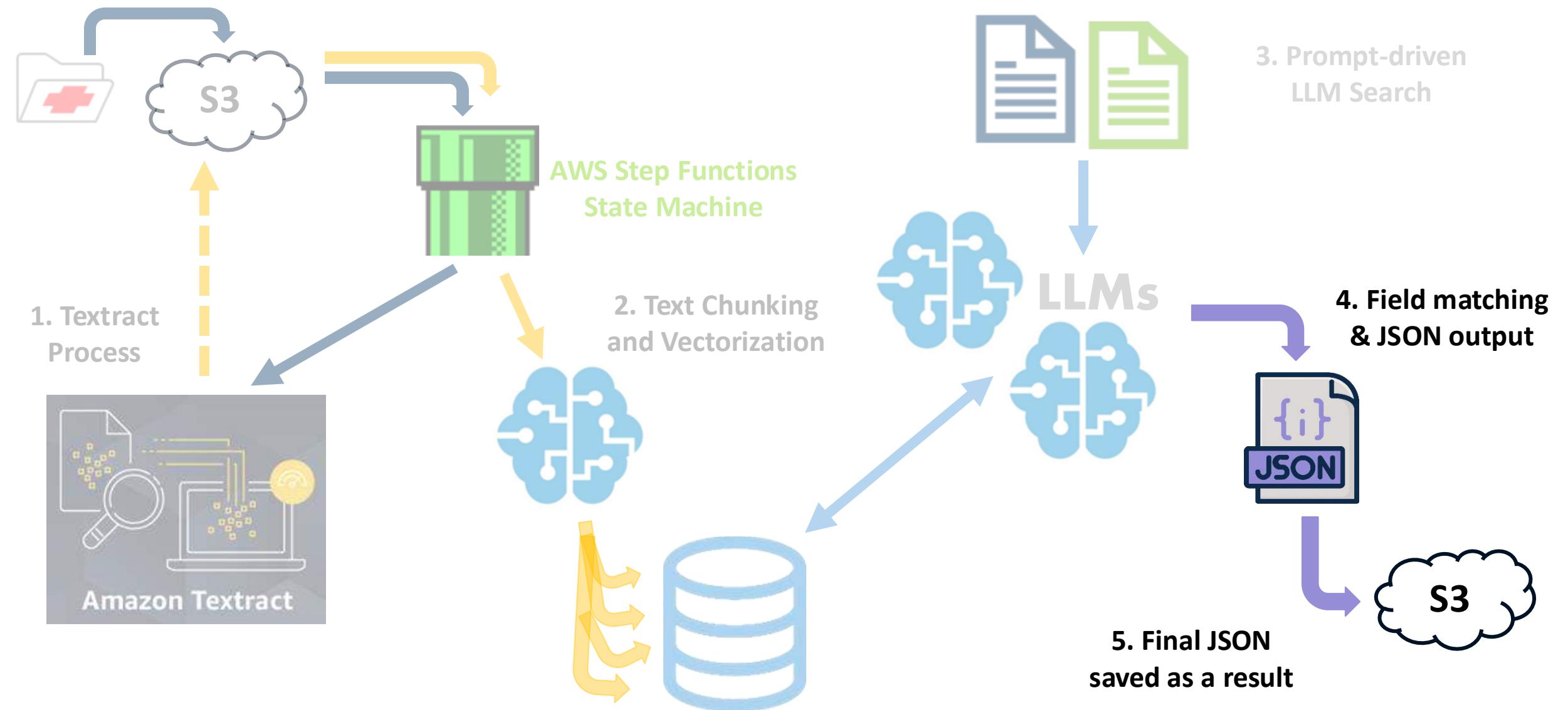
CLAIRE Pilot Project



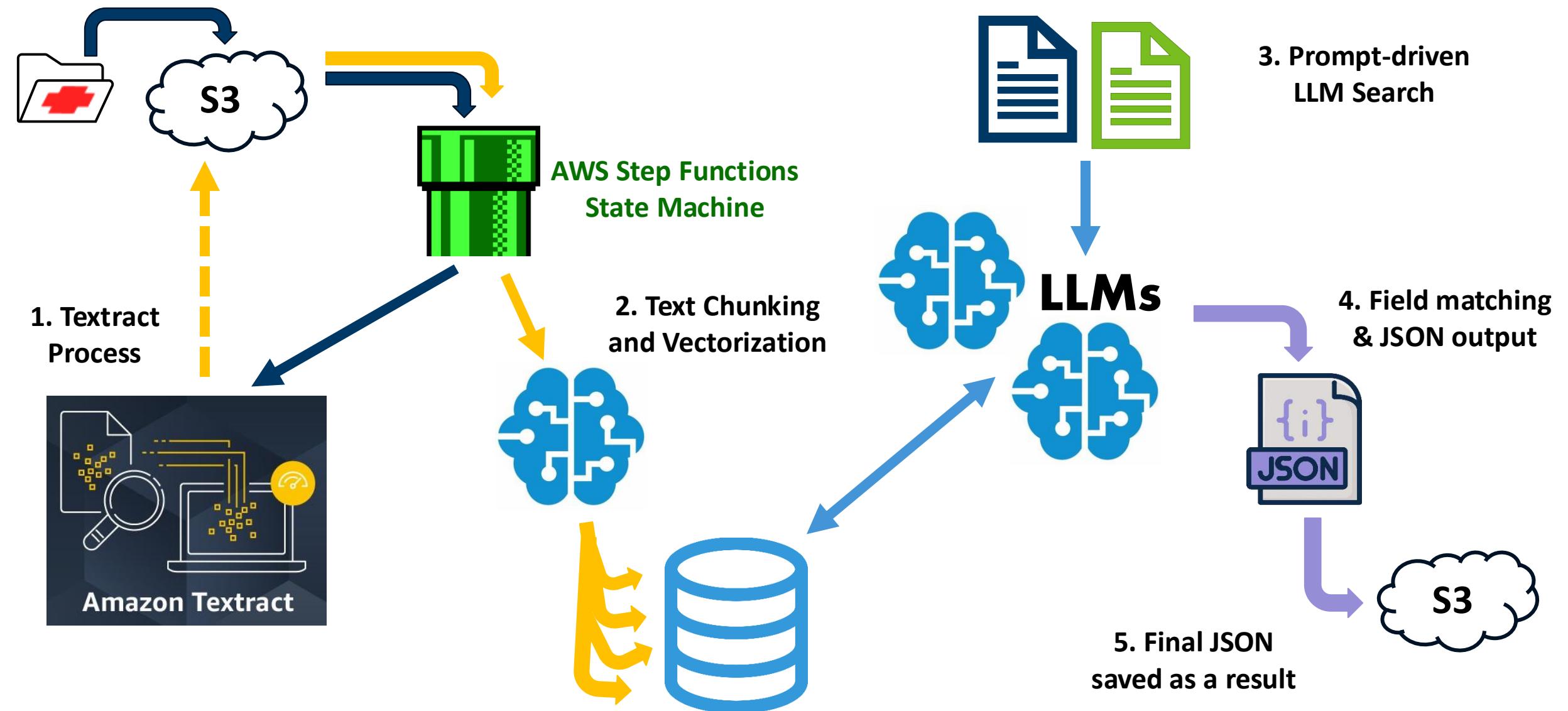
CLAIRE Pilot Project



CLAIRE Pilot Project



CLAIRE Pilot Project



Restructuring the Data

- Throughout the pilot project we looked at various data elements and corresponding accuracy
- We divided data into four categories:
- **STRUCTURED DATA** – data collection better suited to standard code
- **PROMPT DRIVEN SEARCH** – data collection that might require decision-making based upon additional information
- **PROBABILISTIC MATCHING** – data collection that we thought might be best-suited for LLMs
- **SUMMARIZATION** – data collection that might require manual review by staff, but a summary with citations could be helpful and streamline the work

Beneficial Early Results

- We looked at the accuracy by category

Category	Number of variables	Average accuracy across the category
STRUCTURED DATA	76	49.7%
PROMPT DRIVEN SEARCH	50	95.5%
PROBABILISTIC MATCHING	220	92.7%
SUMMARIZATION	33	88.1%

- STRUCTURED DATA variables had the lowest accuracy via large language model assessment
- SUMMARIZATION variables also had a lower accuracy score

Next Steps

- With adequate funding:
 - Continue CLAIRE pilot – more cases, more file types, more providers
 - Other medical chart reviews for other disease areas
 - eCR (Electronic Case Report) use
- Large language models (LLMs) combined with deterministic code may be a smart, hybrid approach to medical chart review, and could result in a more efficient use of staff time
- LLM use could expand our ability to use medical data directly from providers to better understand population health



Thank You!

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Stephanie Meyer

stephanie.meyer@state.mn.us

Public Health Generative AI Use Cases

Agenda – Public Health Use Cases for GenAI

- Electronic Case Reports
- Intelligent Document Processing
- Chatbots
- Generative Business Intelligence and Data Analytics

Data Privacy & Security

Securely build generative AI applications with your data

Data is not used for service improvement



Data remains in the region where API is processed



Data is not shared with model providers



Data is always encrypted in transit and rest



Build with comprehensive data protection & privacy; leverage AWS security services and best practices

Electronic Case Reports

- Problem: Public Health receives thousands of large clinical documents (CDA and HL7) with actionable public health data buried in non-standard data elements, non-standard vocabularies and critical information in free text fields
- Solution: Use Generative AI to ask the questions that public health needs from electronic case reports and extract actionable public health data in a standard format
 - Pregnancy Status (Syphilis in a woman of child bearing age)
 - Occupation (Is the person with Hepatitis A food handler?)

Electronic Case Reports

```
<ClinicalDocument xmlns="urn:hl7-org:v3" xmlns:cda="urn:hl7-org:v3" xmlns:sdtc="urn:hl7-org:sdtc"
  xmlns:voc="http://www.lantanagroup.com/voc" xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
  xsi:schemaLocation="urn:hl7-org:v3 ..../schema/infrastructure/cda/CDA_SDTC.xsd">
  <realmCode code="US"/>
  <typeId extension="POCD_HD000040" root="2.16.840.1.113883.1.3">
    <!-- [C-CDA R1.1] US Realm Header -->
  </typeId>
  <templateId root="2.16.840.1.113883.10.20.22.1.1">
    <!-- [C-CDA R2.1] US Realm Header (V3) -->
  </templateId>
  <templateId extension="2015-08-01" root="2.16.840.1.113883.10.20.22.1.1">
    <!-- [eICR R2 STU1.1] Initial Public Health Case Report Document (eICR) (V2) -->
  </templateId>
  <templateId extension="2016-12-01" root="2.16.840.1.113883.10.20.15.2"/>
  <id root="38e6a983-38ad-484f-a7d2-b294cdbe5435">
    <!-- Globally unique document ID (extension) is scoped by vendor/software -->
  </id>
  <code code="55751-2" codeSystem="2.16.840.1.113883.6.1" codeSystemName="LOINC" displayName="Public Report">
    <!-- Document Code -->
  </code>
  <title>Initial Public Health Case Report</title>
  <effectiveTime value="2020050110515-0400"/>
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  <languageCode code="en-US"/>
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  <versionNumber value="1"/>
  <recordTarget>
    <!-- ***** recordTarget: The patient *****>
    <patientRole>
      <id extension="PT-470127" root="2.16.840.1.113883.19.5">
        <!-- Patient ID -->
      </id>
      <!-- SSN -->
      <id extension="222-22-2222" root="2.16.840.1.113883.4.1"/>
      <addr use="H">
        <streetAddressLine>2222 Home Street</streetAddressLine>
        <city>Sacramento</city>
        <state>CA</state>
        <postalCode>94203</postalCode>
        <county>Sacramento County</county>
        <country>US</country>
      </addr>
      <telecom use="HP" value="tel:555-555-2003"/>
      <telecom use="HP" value="mailto:jose@email.com"/>
    </patientRole>
    <patient>
      <name use="L">
        <given>Joseph</given>
        <family>Patient</family>
      </name>
      <administrativeGenderCode code="M" codeSystem="2.16.840.1.113883.5.1" displayName="Male"/>
      <birthTime value="19890730"/>
      <sdtc:deceasedInd value="false"/>
      <religiousAffiliationCode code="1013" displayName="Christian (non-Catholic, non-specific)" codeSystem="2.16.840.1.113883.5.1076" codeSystemName="HL7 Religious Affiliation"/>
      <raceCode code="1002-5" codeSystem="2.16.840.1.113883.6.238" codeSystemName="Race & Ethnicity" displayName="American Indian or Alaska Native"/>
      <ethnicGroupCode code="2135-2" codeSystem="2.16.840.1.113883.6.238" codeSystemName="Race & Ethnicity - CDC" displayName="Hispanic or Latino"/>
      <languageCommunication>
        <languageCode code="es"/>
        <preferenceInd value="true"/>
      </languageCommunication>
    </patient>
  </recordTarget>
</ClinicalDocument>
```



```
{
  "patient_identifier": "PT-470127",
  "patient_gender": "Male",
  "travel_details": "Recent travel for vacation to Wuhan China from April 22, 2020 to May 5, 2020",
  "medications_received": "Remdesivir 100 mg Injection, 100 mg every day for 4 days",
  "pregnancy_status": "Not specified",
  "symptoms": "Difficulty Breathing (finding)",
  "symptoms_onset_date": "05/13/2020",
  "occupation": "Sales",
  "does_job_involve_food_respond_yes_no_or_unknown": "Unknown",
  "has_chronic_conditions": "Unknown",
  "has_asthma": "Unknown",
  "diagnoses": "Coronavirus as the cause of diseases classified elsewhere (B97.2)",
  "was_hospitalized": "Yes",
  "date_hospitalized": "05/13/2020",
  "date_discharged": "05/13/2020",
  "is_patient_expired": "No",
  "all_medications": "Remdesivir 100 mg Injection",
  "medications_per_condition": "Remdesivir for Coronavirus",
  "eicr_trigger_data": "Coronavirus as the cause of diseases classified elsewhere (B97.2), SARS coronavirus 2 N gene detected",
  "is_this_person_homeless_or_prisoned": "Unknown"
}
```

Intelligent Document Processing

- Problem: Public Health receives paper forms across multiple domains including Vital Records, Electronic Laboratory Reports, Communicable Disease Reports, and WIC applications
- Solution: Use Generative AI to automatically extract the data from paper forms and transform it into formats easily consumable by the systems of record.

Intelligent Document Processing

CDPH
California Department of Public Health

MEASLES (RUBEOLA) CASE REPORT

PATIENT DEMOGRAPHICS

Patient name—last first middle initial Date of birth Age (enter age and check one) Gender
 S P A R E P A N T S S P A R E P A N T S 2 / 1 / 2010 Days Weeks Months Years Male Female

Address—number, street City State ZIP code County
 125 CONCH ST BIKINI BOTTOM CA 93405

Telephone number Home (916) 555 1212 Work () Email: SParépantz@sparepants.com

ETHNICITY (check one) RACE (check all that apply) Asian: Please specify: Pacific Islander: Please specify:
 Hispanic/Latino Black/African-American Asian Indian Himong Thai Native Hawaiian
 Non-Hispanic/ Non-Latino Native American/Alaskan Native Cambodian Japanese Vietnamese Guamanian
 White Chinese Korean Other Asian: Samoan
 Unknown Filipino Laotian Other Pacific Islander:

Country of birth USA Country of residence USA

COMMON LHD TRACKING DATA

CMRID Number IZB Case ID Number WebCMR ID Number
 6342692

Date reported to county 6/12/2025 Date investigation started 6/13/2025 Person/clinician reporting case Patrick Star Reporter telephone ()
 Case investigator completing form Investigator telephone (871) 555 1212 Investigator's jurisdiction

SIGNS AND SYMPTOMS

Rash Rash onset date 6/1/2025 Rash duration 5 days Yes No Unknown Origin on body chest Direction of spread out
 Yes No Unknown Fever Fever onset date 6/1/2025 Was temperature taken Yes Was temperature >=101 (38.3C) If temperature not taken, skin was
 Yes No Unknown Hot Warm Normal Unknown
 Cough Runny nose (coryza) Conjunctivitis Koplik's spots
 Yes No Unknown Yes No Unknown Yes No Unknown
 Other symptoms Describe other symptoms Diagnosis date 6/1/2025
 Yes No Unknown
 Does case meet clinical criteria for further investigation? CASE MEETS CDC/CSTE CLINICAL CRITERIA? (FOR STATE USE ONLY)
 Yes No Unknown
 Yes No Unknown

COMPLICATIONS AND OTHER SYMPTOMS

Hospitalized Days hospitalized Pneumonia Encephalitis Death If yes, date of death
 Yes No Unknown Yes No Unknown Yes No Unknown Yes No Unknown
 Other complications If yes, describe other complications

LABORATORY TESTS

Lab tests done for measles CASE LAB CONFIRMED (FOR LHD USE) CASE LAB CONFIRMED (FOR STATE USE ONLY) LAB RESULT CODES
 Yes No Unknown Yes No Unknown P = Positive
 Serology performed Specimen date Result interpretation N = Negative – Antibody not detected
 Yes No Unknown Yes No Unknown I = Indeterminate
 IgM / / P N I E X U E = Pending
 IgG (acute) / / P N I E X U U N = Not Done
 IgG (convalescent) / / P N I E X U U U = Unknown

Specimen obtained for virus isolation Specimen source Specimen date Virus isolated Name of lab
 Yes No Unknown Nasopharyngeal Urine Other Unknown 6/2/2025 Yes No Unknown

Specimen sent to CDC for genotyping Date sent Virus genotype
 Yes No Unknown / /

Other lab tests performed Other lab test specimen date Specify other lab tests Other lab test results
 Yes No Unknown / /

CDPH 8345 (6/08)

Page 1 of 2

Measles (Rubeola) Case Report—CDPH 8258

VACCINATION/MEDICAL HISTORY

Received one or more doses of measles containing vaccine (MCV) Number of doses
 Yes No Unknown

Dates of vaccination—Dose 1 Dose 2 Dose 3
 / / / / / /

Reason not vaccinated (check all that apply)

1 Personal Belief Exemption (PBE) 4 Lab confirmation of previous disease 7 Delay in starting series or between doses
 2 Permanent Medical Exemption (PME) 5 MD diagnosis of previous disease 8 Other
 3 Temporary Medical Exemption 6 Under age for vaccination 9 Unknown

Prior MD diagnosed measles (see reason 5) Pregnant Yes No Unknown Immunocompromised Yes No Unknown

EPIDEMIOLOGICAL EXPOSURE HISTORY

Spread Setting (check all that apply)

1 Day care 4 Hospital Ward 7 Home 10 College 13 Church
 2 School 5 Hospital ER 8 Work 11 Military 14 International travel
 3 Doctor's office 6 Outpatient hospital clinic 9 Unknown 12 Correctional facility 15 Other

Recent travel or arrival from other country or state within 18 days of rash onset? Yes No Unknown

Countries or states visited Dates in countries or states visited Date of arrival in California
 / / / /

Close contact with person(s) with rash 8-17 days before rash onset? Yes No Unknown

Name	Rash onset date	Relationship	Age (Years)	Same household
1 Sandy Squielle	6/3/2025	friend	13	<input type="checkbox"/> Yes <input checked="" type="checkbox"/> No <input type="checkbox"/> Unknown
2	/ /			<input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown
3	/ /			<input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown

Please list other contacts on a separate sheet or use the contact tracing work sheet.

Epi-linked to a lab-confirmed case? Case name or case ID Outbreak related Outbreak location
 Yes No Unknown Yes No Unknown Bikini Bottom

Import status If case is indigenous, is case If case is imported, describe source
 Indigenous Out-of-state import Imported (linked to imported case) Endemic Unknown Source
 International import Imported virus (viral genetic evidence indicates an imported genotype)

CONTACT INVESTIGATION

Spread Setting (check all that apply)

1 Day care 4 Hospital Ward 7 Home 10 College 13 Church
 2 School 5 Hospital ER 8 Work 11 Military 14 International travel
 3 Doctor's office 6 Outpatient hospital clinic 9 Unknown 12 Correctional facility 15 Other

Number of susceptible contacts Close contacts who have rash 8-17 days after exposure to case (list below)
 Yes No Unknown

Name	Rash onset date	Relationship	Age (Years)	Same household
1 Eugene Krabs	6/4/2025	co-worker	25	<input type="checkbox"/> Yes <input checked="" type="checkbox"/> No <input type="checkbox"/> Unknown
2	/ /			<input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown
3	/ /			<input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown

Please list other contacts on a separate sheet or use the contact tracing work sheet.

CASE CLASSIFICATION (FOR LHD USE) CASE CLASSIFICATION (FOR STATE USE ONLY)

Confirmed Probable Suspect Not a case Unknown Confirmed Probable Suspect Not a case Unknown

MEASLES CASE DEFINITION

Clinical case definition: An illness characterized by all the following: (1) a generalized rash lasting greater than or equal to 3 days, (2) a temperature greater than or equal to 101.0°F (greater than or equal to 38.3°C), and (3) cough, coryza, or conjunctivitis.

Laboratory criteria for diagnosis: Positive serologic test for measles immunoglobulin M antibody; significant rise in measles antibody level by any standard serologic assay; or isolation of measles virus from a clinical specimen.

Case classification
 Suspected: any febrile illness accompanied by rash.
 Probable: a case that meets the clinical case definition, has noncontributory or no serologic or virologic testing, and is not epidemiologically linked to a confirmed case.
 Confirmed: a case that is laboratory confirmed or that meets the clinical case definition and is epidemiologically linked to a confirmed case (a laboratory-confirmed case does not need to meet the clinical case definition).

CDPH 8345 (6/08)

Page 2 of 2

Generative AI Chatbots

- Problem: Public Health departments receive multiple inquiries. Providers using Vital Records and Immunization Information Systems call asking basic questions about how to use these systems. The general public may also call Public Health with simple questions about how to obtain an immunization record or how to enroll in a program like WIC. Although this information is on the Public Health
- Solution: Using Generative AI, resources from public health like user manuals and public facing websites can be the knowledge bases to enable chatbots to automatically answer questions from end users and the general public.

AI Chatbots

Communicable Disease Epi on Call - Guide to Surveillance, Reporting and Control

Interactive Generative AI Assistant

Epi on Call

Customer has joined the chat

BOT 11:25 AM

Welcome to the Self Service Chatbot. How may I assist you with inquiry?

Reference Pages

- [Guide to Surveillance, Reporting and Control](#)
- [Clinical and Laboratory Testing Guidance for Monkeypox](#)

How to Use the Assistant

Our AI documentation assistant is available in the chat window. You can:

- Ask specific questions about the content in the reference pages
- Request information about specific topics
- Get help understanding complex information
- Find relevant sections in the documentation

B I ≡ ≡ ⌂ ☺

What do I do if I find a bat in my room? ➤

End chat **Start a Call**

▼

Generative Business Intelligence

- Problem: Public Health staff do not always have the technical and analytic skills to interact with their data to generate the graphs and stories from their siloed systems to help improve population health.
- Solution: Generative BI enables business users to interact with their data using natural language prompts to build graphs, dashboards and stories that describe their data

AI-assisted storytelling

Interpret data for others

Help others derive meaning from data and reach conclusions to drive decisions

Generate stories using AI

Produce cohesive, powerful, and insightful narratives

Create refined content

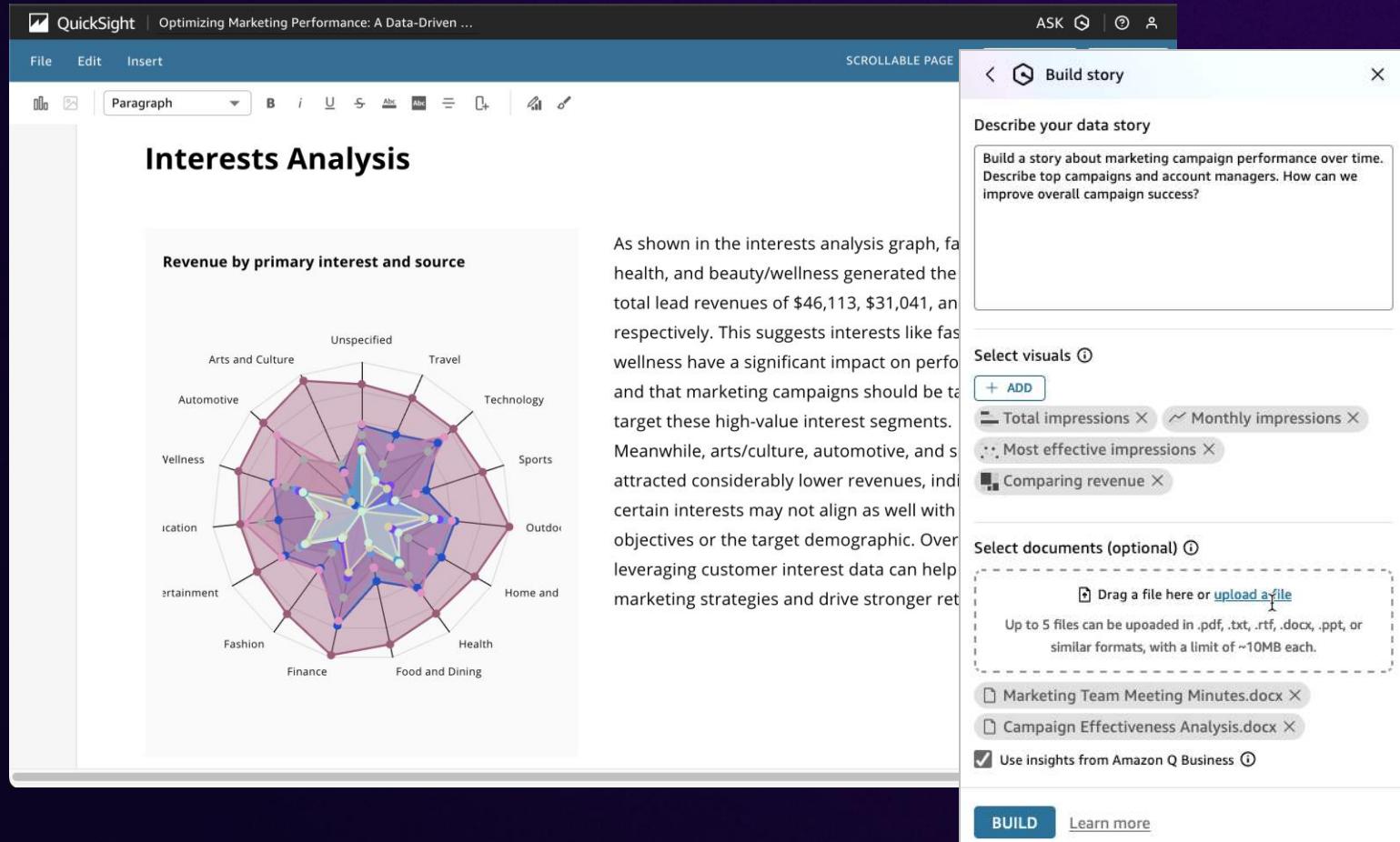
Control AI verbosity, customize narrative text, and apply stunning visual themes to bring content to life

Augment with unstructured data

Upload files with related content to enrich the story with additional insights

Share up-to-date governed data

Quickly update and disseminate data at any time



Generative Business Intelligence

Build visuals

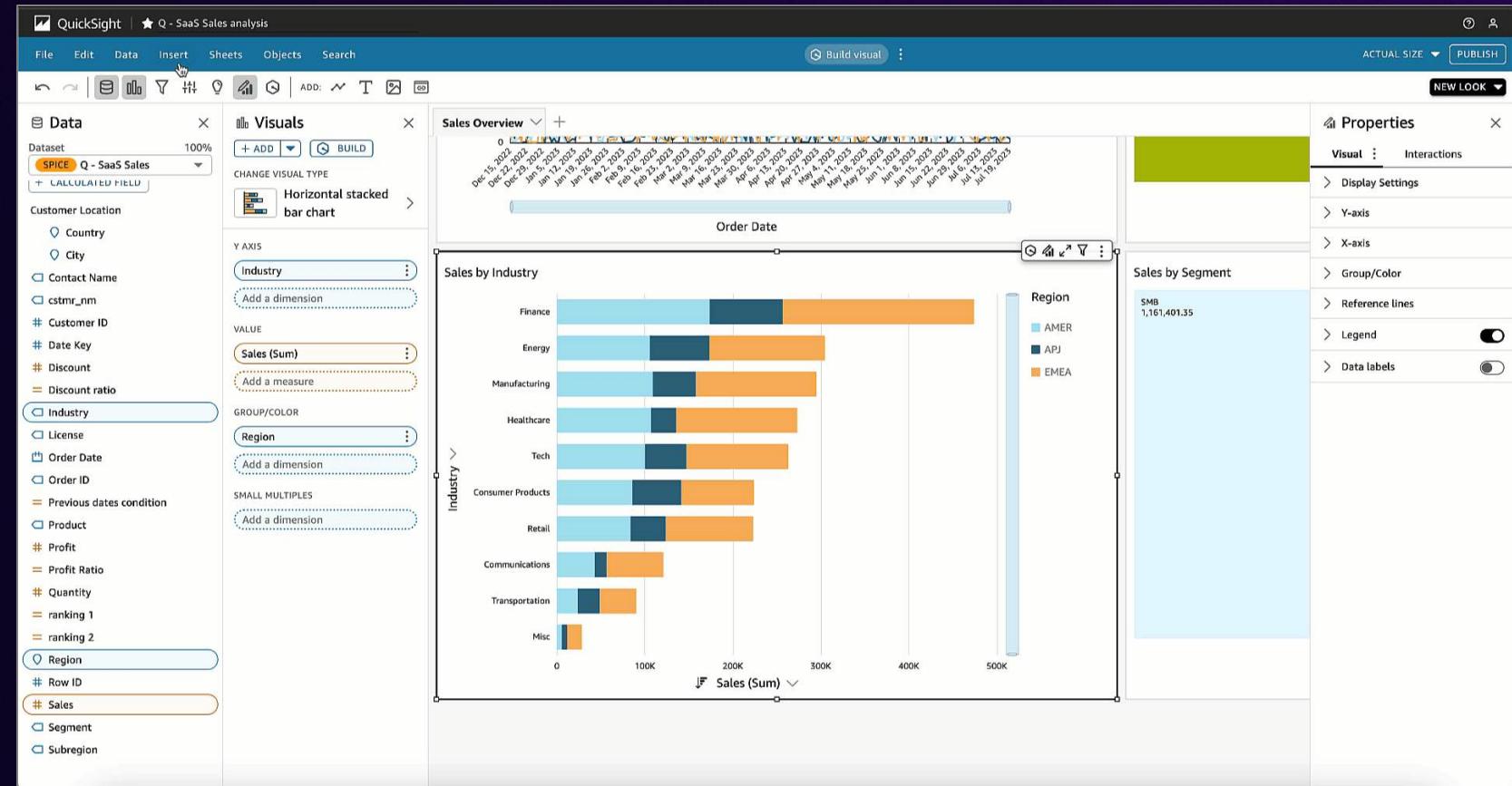
Use natural language to quickly build visuals for dashboards and reports

Build calculations

Easily create calculations using natural language without looking up or learning specific syntax

Refine visuals

Quickly update visuals by describing desired formats using natural language





Blog: Transforming electronic case reporting with generative AI: Unlocking faster public health responses



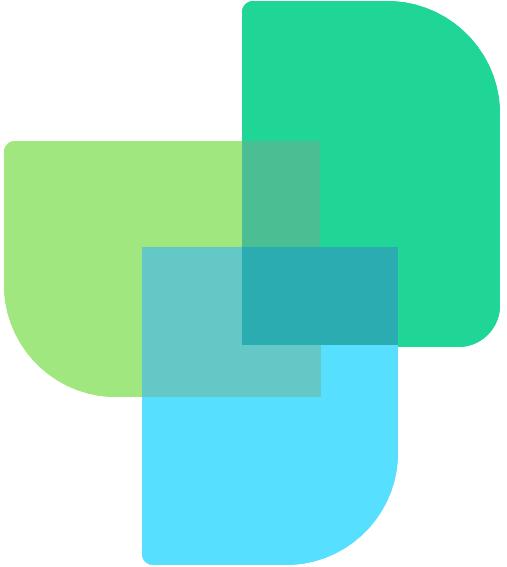
Blog: Scalable intelligent document processing using Amazon Bedrock



Thank you!



Audience Q & A



Thank You!

UP NEXT...

5-6pm Central Time | Join us for our *Data Modernization in Motion Networking Reception* – Park View, 4th Floor

TOMORROW...

8:30-9:30am Central Time | Hashbrowns & High Tech Peer Networking Breakfast

9:45am Central Time | Concurrent sessions continue, followed by preconference closing plenary