



# ATLAS Deepdive: Cohorts and Concept Sets



**OHDSI Community Call**  
**June 17, 2025 • 11 am ET**



# Upcoming Community Calls

Date	Topic
June 17	ATLAS Deepdive: Cohorts and Conceptsets
June 24	ATLAS Deepdive: Characterization, Cohort Pathways, Incidence
July 1	ATLAS Deepdive: Technical and Administrative Capabilities
July 8	No Meeting – Europe Symposium
July 15	Europe Symposium Review
July 22	OMOP/OHDSI Research Spotlight
July 29	Asia-Pacific Regional Updates
Aug. 5	No Meeting
Aug. 12	Newcomer Introductions



# Three Stages of The Journey

**Where Have We Been?**

**Where Are We Now?**

**Where Are We Going?**





# OHDSI Shoutouts!



Congratulations to the team of  
**Meghan L. McCarthy, Jonah  
Bradenday, Elizabeth Chen, Mark  
R. Zonfrillo, and Indra Neil Sarkar**  
on the publication of **Reductions in  
Blood Lead Level Screening During  
Peak COVID-19 Restrictions and  
Beyond** in *Public Health  
Challenges*.

Public Health Challenges

WILEY

Public Health CHALLENGES  
Open Access

RESEARCH ARTICLE OPEN ACCESS

## Reductions in Blood Lead Level Screening During Peak COVID-19 Restrictions and Beyond

Meghan L. McCarthy<sup>1</sup> | Jonah Bradenday<sup>2</sup> | Elizabeth Chen<sup>1,2</sup> | Mark R. Zonfrillo<sup>3,4</sup> | Indra Neil Sarkar<sup>1,2,5</sup>

<sup>1</sup>Warren Alpert Medical School of Brown University, Providence, Rhode Island, USA | <sup>2</sup>Brown Center for Biomedical Informatics, Providence, Rhode Island, USA | <sup>3</sup>Department of Emergency Medicine, Warren Alpert Medical School of Brown University, Providence, Rhode Island, USA | <sup>4</sup>Department of Pediatrics, Warren Alpert Medical School of Brown University, Providence, Rhode Island, USA | <sup>5</sup>Rhode Island Quality Institute, Providence, Rhode Island, USA

Correspondence: Indra Neil Sarkar ([neil\\_sarkar@brown.edu](mailto:neil_sarkar@brown.edu))

Received: 14 May 2024 | Revised: 21 October 2024 | Accepted: 7 December 2024

**Funding:** The project described was supported by an Institutional Development Award for Clinical and Translational Research (U54GM115677) from the National Institute of General Medical Sciences of the National Institutes of Health, which funds Advance Rhode Island Clinical and Translational Research (Advance RI-CTR). This project was also supported by National Institutes of Health (U54GM104942), as well as by the Warren Alpert Medical School of Brown University's Primary Care and Population Medicine Summer Research Funding program.

**Keywords:** health information exchange | lead exposure | lead screening | pediatric primary care | screening | well child care

### ABSTRACT

**Background and Objectives:** Among the multitude of health effects on children associated with the COVID-19 pandemic, there have been significant interruptions in the provision of routine pediatric primary care, including blood lead level (BLL) screening. We aimed to investigate trends in BLL screening before and during the pandemic era using patient-level electronic health record data extracted from CurrentCare, Rhode Island's statewide health information exchange (HIE).

**Methods:** De-identified data were analyzed from CurrentCare for the study period January 2018 to December 2021. We utilized ATLAS, a web-based analytics platform from the Observational Health Data Sciences and Informatics (OHDSI) community, to extract and stratify BLL by variables of interest from the CurrentCare data, standardized to OHDSI's Observational Medical Outcomes Partnership common data model.

**Results:** A decrease in BLL screening occurred in the spring of 2020, aligning with initial periods of shelter-in-place in response to the novel coronavirus outbreak; there was a 48% decrease comparing quarter 2 (April to June) of 2019 and 2020. BLL screening rebounded in the summer of 2020, however, it remained 16% lower overall in 2020 than in 2019. In 2021, BLL screening fell again to 23% lower than in 2019. Although overall numbers of BLL screenings were reduced, the proportion of abnormal BLLs was higher, particularly in the range of 3.5–5.0 µg/dL.

**Conclusions:** Leveraging statewide HIE data, we found that significant deficiencies in BLL screening remain unresolved since the beginning of the COVID-19 pandemic. The disruption of children's lives by the COVID-19 pandemic appears to have greatly affected lead screening and exposure in Rhode Island.



# OHDSI Shoutouts!



Congratulations to the team of **Zhang Meng, Shen Peng, Liu Zhike, Mui Van Zandt, Li Jing, Li Chao, Sun Yexiang, Xie Junqing, Wan Eric Yuk Fai, George Hripcsak, Chen Yong, Lin Hongbo, Zhan Siyan, and Sun Feng** on the publication of **Study of application of Common Data Model of Observational Medical Outcomes Partnership in China** in the *Chinese Journal of Epidemiology*.

## Study of application of Common Data Model of Observational Medical Outcomes Partnership in China

Zhang Meng · Shen Peng · Liu Zhike · Van Zandt Mui · Li Jing · Li Chao · Sun Yexiang · Xie Junqing · Wan Eric Yuk Fai · George Hripcsak · Chen Yong · Lin Hongbo

Zhan Siyan · Sun Feng

[Authors Info & Affiliations](#)

Published : 2025-05-10 · DOI: [10.3760/cma.j.cn112338-20240924-00595](https://doi.org/10.3760/cma.j.cn112338-20240924-00595)

PDF Download

8 4 0 0 0 0 0

PDF Download

### summary

**Objective** To summarize the application status of the Observational Medical Outcomes Partnership (OMOP) Common Data Model (CDM) in China and provide a reference for the implementation strategy of data standardization and evidence sharing in China.

**Methods** PubMed, Embase, Web of Science, CNKI, VIP, Wanfang Data Knowledge Service Platform, and Chinese Biomedical Literature System were searched for relevant studies on Chinese medical institutions/platforms using OMOP CDM to standardize data since their inception until March 15, 2023. Information such as institutions, patient types, and numbers were extracted.

### Cite this article

### PERMISSIONS

Request permissions for this article from CCC.

REQUEST PERMISSIONS



# OHDSI Shoutouts!



Congratulations to the team of  
**Jacob Zelko and Justin  
Manjourides** on the publication  
of **A Generalized Tool to Assess  
Algorithmic Fairness in Disease  
Phenotype Definitions in the  
Proceedings — AMIA Joint  
Summits on Translational Science.**

## A Generalized Tool to Assess Algorithmic Fairness in Disease Phenotype Definitions

Jacob S. Zelko, B.S.<sup>1</sup>, Justin Manjourides, Ph.D.<sup>2, 3</sup>

<sup>1</sup>Department of Mathematics, Northeastern University, Boston, MA, USA;

<sup>2</sup>Department of Public Health and Health Sciences, Northeastern University, Boston, MA, USA;

<sup>3</sup>Roux Institute, Northeastern University, Portland, ME, USA

### Abstract

*For evidence from observational studies to be reliable, researchers must ensure that the patient populations of interest are accurately defined. However, disease definitions can be extremely difficult to standardize and implement accurately across different datasets and study requirements. Furthermore, in this context, they must also ensure that populations are represented fairly to accurately reflect populations' various demographic dynamics and to not overgeneralize across non-applicable populations. In this work, we present a generalized tool to assess the fairness of disease definitions by evaluating their implementation across common fairness metrics. Our approach calculates fairness metrics and provides a robust method to examine coarse and strongly intersecting populations across many characteristics. We highlight workflows when working with disease definitions, provide an example analysis using an OMOP CDM patient database<sup>1</sup>, and discuss potential directions for future improvement and research.*

### Introduction

Observational health research uses data collected by healthcare providers (e.g. hospital systems, clinics, etc.) to conduct retrospective population level analyses of various patient populations. [2] This data, commonly referred to as "Real World Data" [3], are data which relates to patient health status and/or the delivery of health care that is routinely collected from a variety of sources including electronic health records, medical claims and billing activities, and product and disease registries. Evidence generated from analyses using "Real-world Data" (RWD) from these types of sources is referred to as "Real-world Evidence" (RWE) and can be used to evaluate the effectiveness and safety of medical treatments, interventions, and healthcare practices in everyday clinical settings, outside of controlled clinical trials. [3]

Within observational health research, several communities of practice have emerged dedicated to how best to make use of this data. [4, 5, 6] One community that has emerged over the past decade is the the Observational Health Data Sciences and Informatics (OHDSI) open science collaborative. [7] This collective consists of hundreds of researchers with expertise in clinical sciences, health informatics, healthcare economics, and many more domains where observational health has found application. One technology that emerged from the OHDSI collaborative is the Observational Medical Outcomes Partnership Common Data Model (OMOP CDM)<sup>2</sup>.



# OHDSI Shoutouts!



Congratulations to the team of **Seok Kim, Dachung Boo, Sooyoung Yoo, Borham Kim, Kyubo Kim, Kwangsoo Kim, Eunhye Song, Junmo Kim, Hyun Gee Ryoo, Jin Chul Paeng, In Young Choi, SooJeong Ko, le Ryung Yoo, Rae Woong Park, and Ho-Young Lee** on the publication of **Secondary Cancer Risk in Breast Cancer with and without Radiotherapy: The Observational Health Data Sciences and Informatics (OHDSI) Cohort Study** in *Cancer Research and Treatment*.







# OHDSI Shoutouts!



Congratulations to the team of **Kim López-Güell, Martí Català, Daniel Dedman, Talita Duarte-Salles, Raivo Kolde, Raúl López-Blasco, Álvaro Martínez, Gregoire Mercier, Alicia Abellan, Johnmary T. Arinze, Theresa Burkard, Edward Burn, Zara Cuccu, Antonella Delmestri, Dominique Delseny, Sara Khalid, Chungsoo Kim, Ji-woo Kim, Kristin Kostka, Cora Loste, Miguel A. Mayer, Jaime Meléndez-Cardiel, Núria Mercadé-Besora, Mees Mosseveld, Akihito Nishimura, Hedvig ME. Nordeng, Jessie O. Oyinlola, Roger Paredes, Laura Pérez-Crespo, Marta Pineda-Moncusí, Juan Manuel Ramírez-Anguita, Nhung TH. Trinh, Anneli Uusküla, Bernardo Valdivieso, Daniel Prieto-Alhambra, Junqing Xie, Lourdes Mateu, and Annika M. Jödicke** on the publication of **Clusters of post-acute COVID-19 symptoms: a latent class analysis across 9 databases and 7 countries** in the *Journal of Clinical Epidemiology*.



Journal of Clinical Epidemiology

Available online 13 June 2025, 111867

[In Press, Journal Pre-proof](#) [What's this?](#)



Original Research

## Clusters of post-acute COVID-19 symptoms: a latent class analysis across 9 databases and 7 countries

Kim López-Güell<sup>1 #</sup>, Martí Català<sup>1 #</sup>, Daniel Dedman<sup>2</sup>, Talita Duarte-Salles<sup>3 4</sup>, Raivo Kolde<sup>5</sup>, Raúl López-Blasco<sup>6</sup>, Álvaro Martínez<sup>7</sup>, Gregoire Mercier<sup>9 10</sup>, Alicia Abellan<sup>3</sup>, Johnmary T. Arinze<sup>4</sup>, Theresa Burkard<sup>1</sup>, Edward Burn<sup>1</sup>, Zara Cuccu<sup>2</sup>, Antonella Delmestri<sup>1</sup>, Dominique Delseny<sup>9</sup>, Sara Khalid<sup>1</sup>, Chungsoo Kim<sup>11</sup>, Ji-woo Kim<sup>12</sup>, Kristin Kostka<sup>1 13</sup>, Cora Loste<sup>8 21 23</sup>...  
Annika M. Jödicke<sup>1 †</sup>

[Show more](#) ▾

[+](#) Add to Mendeley [🔗](#) Share [📄](#) Cite

<https://doi.org/10.1016/j.jclinepi.2025.111867>

[Get rights and content](#) ↗

Under a Creative Commons [license](#) ↗

[Open access](#)

### Highlights

- 787,078 persons with PCC were included, which makes this study the largest international PCC study to our knowledge
- Complex multi-symptomatic clusters included anxiety-depression, abdominal-gastrointestinal symptoms, and respiratory problems with fatigue and joint pain





# Three Stages of The Journey

**Where Have We Been?**

**Where Are We Now?**

**Where Are We Going?**





# Upcoming Workgroup Calls



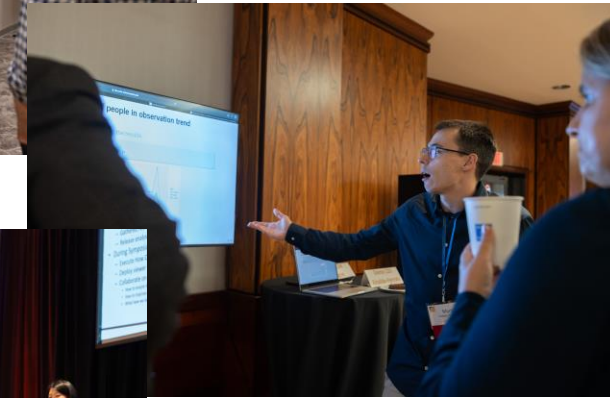
Date	Time (ET)	Meeting
Tuesday	12 pm	ATLAS
Thursday	8 am	India Community Call
Thursday	9 am	Oncology Vocabulary/Development Subgroup
Thursday	11 am	Themis
Thursday	12 pm	HADES
Friday	10 am	GIS-Geographic Information System
Friday	11:30 am	Steering
Monday	9 am	Vaccine Vocabulary
Monday	10 am	Africa Chapter
Monday	10 am	Getting Started Subgroup
Tuesday	9 am	Oncology Genomic Subgroup
Tuesday	9:30 am	CDM Survey Subgroup



# TWO Weeks Remaining

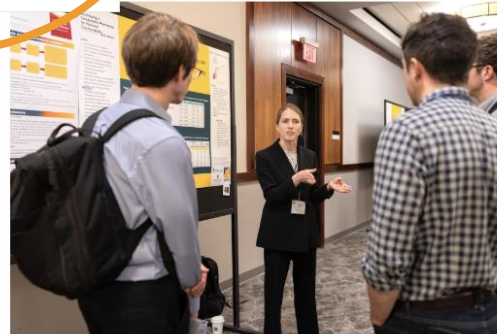
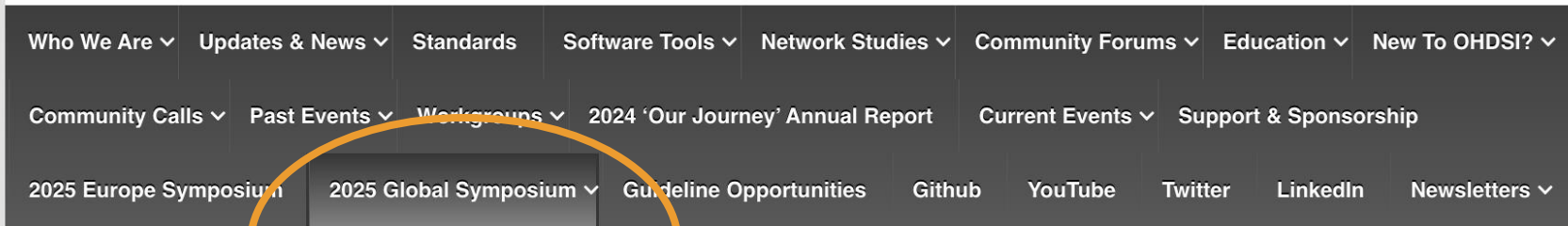
The submission deadline for the 2025 Global Symposium Collaborator Showcase is **July 1.**

More information about the collaborator showcase, including links to the submission form and poster templates, can be found on the #OHDSI2025 homepage.





# Global Symposium: Oct. 7-9



## 2025 OHDSI Global Symposium

Oct. 7-9 • New Brunswick, N.J. • Hyatt Regency Hotel

There is nothing quite like the OHDSI Global Symposium, which welcomes hundreds of collaborators around the world who believe in the shared mission of improving health by empowering a community to collaboratively generate the evidence that promotes better health decisions and better care. We can't wait to return for our biggest event of the year this October in New Brunswick, N.J.





# Europe Symposium Agenda

## Symposium Agenda – July 7, 2025

Time	Topic
8:00 – 9:00	Registration & Coffee
9:00 – 9:10	Welcome to the European OHDSI Journey ( <i>Speakers: Liesbet M. Peeters &amp; Peter Rijnbeek</i> )
9:10 – 9:30	Journey of OHDSI: Where have we been and where can we go together? ( <i>Speaker: Patrick Ryan</i> )
9:30 – 11:00	Impact of Leveraging OMOP CDM for Scalable and Reliable Evidence Generation Showcased by the National Nodes ( <i>Moderators: Renske Los &amp; Annelies Verbiest</i> )
11:00 – 11:30	Coffee Break
11:30 – 12:45	Collaborator Showcase: Rapid Fire Presentations ( <i>Moderator: TBC</i> )
12:45 – 13:45	Lunch
13:45 – 16:00	OHDSI Collaborator Showcase Early Investigator Mentor Meeting (14:00 – 15:00)
16:00 – 17:10	Bridging Policy and Practice: OHDSI's Role in Implementing the European Health Data Space ( <i>Panel debate</i> ) ( <i>Confirmed speakers/moderators: Enrique Bernal-Delgado, Nick Marly, Talita Duarte-Salles, Patrick Ryan, Dipak Kalra</i> )
17:10 – 17:30	Closing remarks ( <i>Speakers: Liesbet M. Peeters &amp; Peter Rijnbeek</i> )

## Agenda Saturday July 5, 2025

Time	Activity	Track 1A – Newcomers	Track 1B – Newcomers	Track 2 – Advanced	Track 3 – NN/WG
09:30 – 10:00		Registration + coffee			
10:00 – 12:30	Morning Session	<b>Introduction to OHDSI – Tutorial</b> Lead: Renske Los, Aniek Markus & Laura Verbeij (Erasmus MC) Overview of OHDSI, key concepts, and an introduction to the OMOP Common Data Model			<b>HADES hack-a-thon</b> Lead: Martijn Schuermie (J&J), Adam Black (Erasmus MC), Anthony Sena (Janssen R&D)  Hands-on coding and tool development in HADES
12:30 – 13:30		Lunch break			
13:30 – 15:00	Afternoon Session I	<b>OMOP CDM &amp; ETL Conventions</b> Lead: Maxim Mainat (Erasmus MC), Sofia Bazakou & Anne van Winzum (The Hyve)	<b>OHDSI Standardized Vocabularies for Research – Part 1.1</b> Lead: Anna Ostropelets (Janssen R&D), Polina Talapova (Sciforce), Vlad Korsik & Oleg Zhuk (Odysseus)  Concept sets & patient identification techniques.		
15:00 – 15:30		Coffee Break			
15:30 – 17:00	Afternoon Session II		<b>OHDSI Standardized Vocabularies for Research – Part 1.2</b> Lead: Anna Ostropelets (Janssen R&D), Polina Talapova (Sciforce), Vlad Korsik & Oleg Zhuk (Odysseus)  Concept sets & patient identification techniques.		
17:15 – 18:45*		*Optional – guided city tour Hasselt (with local specialties)			

## Agenda Sunday July 6, 2025

Time	Activity	Track 1A – Newcomers	Track 1B – Newcomers	Track 2 – Advanced	Track 3 – NN/WG
09:30 – 10:00		Registration + coffee			
10:00 – 12:30	Morning Session		<b>OHDSI Standardized Vocabularies for Research – Part 2</b> Lead: Anna Ostropelets (Janssen R&D), Polina Talapova (Sciforce), Vlad Korsik & Oleg Zhuk (Odysseus)  Final discussion & application of concept sets.	<b>NN All Actors Meet Parallel NN meetings</b>	
12:30 – 13:30		<b>Data Partners Lunch Break</b>			
13:30 – 15:00	Afternoon Session I	<b>Whirlwind introduction to Open-Source Analytic Tools – Part 1</b> Lead: Martijn Schuermie (J&J), Adam Black (Erasmus MC), Anthony Sena (Janssen R&D) Overview of HADES and other key OHDSI tools for analysis.		<b>Running characterisation studies from beginning to end: a tutorial using DARWIN EU standardised analytics – Part 1</b> Lead: Daniel Prieto-Alhambra (Oxford University)	<b>NN All Actors Meet Parallel NN meetings</b>
15:00 – 15:30		<b>Coffee Break</b>			
15:30 – 17:00	Afternoon Session II	<b>Whirlwind introduction to Open-Source Analytic Tools – Part 2</b> Lead: Martijn Schuermie (J&J), Adam Black (Erasmus MC), Anthony Sena (Janssen R&D) Overview of HADES and other key OHDSI tools for analysis.		<b>Running characterisation studies from beginning to end: a tutorial using DARWIN EU standardised analytics – Part 2</b> Lead: Daniel Prieto-Alhambra (Oxford University)	<b>OHDSI Europe NN leads meet</b> Lead: Renske Los (only NN leads/managers)
17:00 – 18:00*		<b>*Optional – networking drink</b>			



# Columbia Summer School on OHDSI

Registration is open for the first ever Columbia Summer School on OHDSI, held July 14-18, 2025, at the Columbia University Department of Biomedical Informatics in New York City.

The Columbia Summer School in Observational Health Data Science and Informatics, Artificial Intelligence, and Real World Evidence (RWE) offers health professionals, researchers and industry practitioners the opportunity to gain familiarity and hands-on experience with real world data and generating real world evidence. Participants will learn about the different types of healthcare data captured during routine clinical care, including electronic health records and administrative records, and how these data can be standardized to the OMOP Common Data Model to enable distributed data network research.



## Meet Our Faculty



**George Hripcsak, MD MS**  
Vivian Beaumont Allen  
Professor of Biomedical  
Informatics



**Patrick Ryan, PhD**  
Adjunct Assistant  
Professor of Biomedical  
Informatics



**Anna Ostropolets, MD PhD**  
Adjunct Assistant  
Professor of Biomedical  
Informatics



**Karthik Natarajan, PhD**  
Assistant Professor of  
Biomedical Informatics





# OHDSI on Bluesky

**OHDSI is now on Bluesky!**

**You can now get updates on all community activities and see all global research through the #OHDSISocialShowcase on Bluesky.**



[bsky.app/profile/ohdsi.bsky.social](https://bsky.app/profile/ohdsi.bsky.social)



# Prediction of Hyponatremia in Cancer Patients Using Machine Learning Based on Oncology CDM

(Yeji Lee, Hyunwoo Park, Yul Hwangbo, HyoSung Cha)



**OHDS**

<sup>3</sup> Graduate School of Cancer Science and Policy, National Cancer Center, Goyang, South Korea

The number of cancer patients worldwide has been steadily increasing. Although the survival rate of cancer patients has improved due to advances in treatment technologies and preventive policies, the management of complications in cancer survivors remains a critical concern. Electrolyte abnormalities are common among cancer patients but are often overlooked. Mild electrolyte imbalances often do not present symptoms and are frequently detected through routine laboratory tests. However, if it severe, it can cause serious complications. In particular, hyponatremia is a common complication in cancer patients and requires early detection, because it is associated with increased long-term hospitalization and mortality. This study developed a previously unexplored model for predicting the occurrence of hyponatremia in cancer patients based on the Oncology Common Data Model (CDM).

**Data source:**

**Data source:**

The oncology CDM (version 5.4), converted from data of 119,854 cancer patients who visited the National Cancer Center from January 2010 to December 2021, was used in the analysis.

**Cohort Selection Criteria:**

- Exclusion criteria:**
- (1) patients who were not diagnosed with cancer before the onset of hyponatremia were excluded.
  - (2) Patients diagnosed with liver cancer or liver disease were excluded because hyponatremia could be induced due to decreased liver function.
  - (3) Patients who developed hyponatremia 365 days after cancer diagnosis were excluded.
  - (4) To develop a predictive model one day before the onset of hyponatremia, the observation period was defined from the date of cancer diagnosis to the onset of hyponatremia.

Finally, 33,476 patients were included in the analysis.

- Hyponatremia group: 3,102 patients (9.3%)

- Normonatremia group: 30,374 patients (90.7%)

**Outcome:**

Occurrence of hyponatremia (serum sodium  $\leq 125$  mmol/L).

**Model Development:**

We developed a stacking ensemble model using logistic regression(LR), random forest(RF), support vector machine(SVM), and light gradient boosting(LGB) as a base model, and extreme gradient boosting(XGB) as a meta model.

### Model Validation and Evaluation:

The dataset was divided into training(80%) and test(20%) sets, and 5-fold cross-validation was applied to the training set. The performance of the model was evaluated with the base model and stacking ensemble using area under the receiver operating characteristic curve(AUROC), accuracy, sensitivity, specificity, positive predictive value(PPV), negative predictive value(NPV), and F1 scores. The Shapley Additive Explanations(SHAP) algorithm was used to visually express the effect of each variable on the prediction of hyponatremia.

## Results

Most models demonstrated excellent performance in AUROC, accuracy, and F1 score, but relatively low sensitivity and Positive Predictive Value (PPV). (Table 1). To address these limitations, we employed a stacking ensemble model, which achieved consistently high performance across all evaluation metrics.

Table 1. Model Performance of Hyponatremia Prediction

Models	AUR	Accuracy	Sensitivity	Specificity	PPV	NPV	F1
LR	0.899	0.929	0.487	0.976	0.683	0.947	0.765
RF	0.899	0.919	0.184	0.996	0.837	0.920	0.629
<b>SVM</b>	0.899	0.925	0.295	0.992	0.787	0.930	0.695
LGB	0.899	0.937	0.506	0.982	0.752	0.950	0.785
XGB	0.930	0.938	0.511	0.984	0.768	0.950	0.614
<b>Stack Ens</b>	<b>0.922</b>	<b>0.939</b>	<b>0.753</b>	<b>0.969</b>	<b>0.855</b>	<b>0.969</b>	<b>0.793</b>

The SHAP summary plot shows the relative importance of each clinical feature in predicting hyponatremia. (Figure 2). The higher the age and the lower the chloride level, the greater the likelihood of developing hyponatremia. This is consistent with previous research findings, as sodium and chloride levels are closely correlated. Cancer-specific information found that SEER stage 7 and M stage 1 were associated with a high incidence of hyponatremia, indicating a high probability of hyponatremia in patients with tumor metastasis.

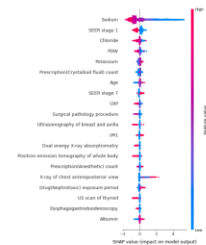


Figure 2. SHAP summary plot of the LightGBM model

## Conclusions

In this study, we developed a predictive model to forecast the occurrence of hypotension one day in advance in cancer patients undergoing the oncology CDM. Laboratory tests, age, and progression of cancer were identified as major predictors. A review of recent sodium test findings and test dates before hypotension confirmed a sharp drop in sodium levels above 15 mmol/L without a sodium test for more than 100 days in some patients. Based on the major predictors identified in this study, providing baseline data for the early detection of hypotension enables medical staff to anticipate and prevent its occurrence in advance. Furthermore, patients requiring sodium monitoring can be identified proactively, thereby reducing the risk of overlooking patients for extended periods. However, as this study was based on a single center, the generalizability of the model needs to be confirmed in the multicenter data of various patients. We are currently exploring various federated learning platforms and are considering FederNet, which is actively utilized in large hospitals.

This study was supported by the Bio-Industrial Technology Development Program (20014841) funded By the Ministry of Trade, Industry & Energy (MOTIE, Korea).

Contact: leeyeji14@ncc.re.kr



**www.ohdsi.org**

## #JoinTheJourney





# #OHDSISocialShowcase This Week

## Tuesday

# Trend analysis in Prevalence of Dementia Medications: a perspective from Taipei Medical University

(**Septi Melisa**, Phan Thanh-Phuc, Nguyen Phung-Anh, Jason C. Hsu)



The utilization of dementia medications has significantly increased over the observed period, with a particularly notable surge occurring between 2016 and 2020

### *Trend analysis in Prevalence of Dementia Medications: a perspective from Taipei Medical University*

PRESENTER: **Septi Melisa**

d931111003@tmu.edu.tw

**Introduction:** The global prevalence of diagnosed dementia is expected to increase dramatically from 57.4 million cases in 2019, with an uncertainty range of 50.4-65.1 million, to nearly three times this figure by 2050. Further research is required to gain a deeper understanding of the utilization, distribution, and prescribing trends of dementia medications.

### Methods

**Data source:** Taipei Medical University Clinical Research Database (TMUCRD) which was mapped into OMOP-CDM

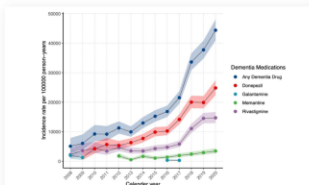
**Target Population:** Patients who have the condition occurrence of dementia for the first time in the patient's history, with continuous observation of at least 0 days after event index date and limit initial events to earliest event per person. Subjects should also receive medication for dementia (Donepezil, Rivastigmine, Galantamine, and Memantine) to ensure that they meet the accurate diagnosis.

**Analysis:** The analysis was conducted using IncidencePrevalence package version 0.7.4

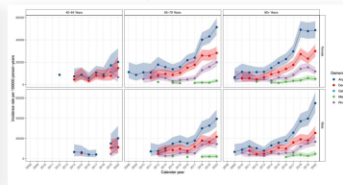
Diagnosis	ICD 10	SNOMED-CT
Dementia	F02.x, F03.x, G31.8	52448006, 56267009, 15662003, 230270009, 12348006, 230288001, 762707000, 73941000119107
Alzheimer	F00.x G30.x	28929004
Vascular Dementia	F01.x	429998004
ATC Name	ATC Code	RxNorm Ingredient Code
Donepezil	N06DA02	135447
Rivastigmine	N06DA03	183379
Galantamine	N06DA04	4637
Memantine	N06DX01	6719

### Results

**Figure 1:** Trends in Dementia Medication Incidence Rates (2008–2020)



**Figure 2:** Incidence Rates of Dementia Medication Use by Age Group and Gender (2008–2020)



**Conclusion:** Donepezil remains the most frequently prescribed drug for dementia, followed by Rivastigmine, both demonstrating significant growth in usage. In contrast, Galantamine and Memantine show relatively lower and more stable usage patterns. A stratified analysis by age and gender reveals that the highest prescription rates occur in the oldest age group (80+ years), with women generally exhibiting higher usage rates than men. This trend underscores an increasing reliance on dementia medications, particularly among older adults, signalling a rising demand for dementia care in this demographic.



▲ **Septi Melisa<sup>1</sup>; Phan Thanh-Phuc<sup>1</sup>;  
Nguyen Phung-Anh<sup>2,3,4</sup>; Jason C. Hsu<sup>1,2,3,4\*</sup>**

<sup>1</sup> International Ph.D. program in Biotech and Healthcare Management, College of Management, Taipei Medical University, Taipei, Taiwan  
<sup>2</sup> Clinical Data Center, Office of Data Science, Taipei Medical University, Taipei, Taiwan  
<sup>3</sup> Research Center of Health Care Industry Data Science, College of Management, Taipei Medical University, Taipei, Taiwan  
<sup>4</sup> Clinical Big Data Research Center, Taipei Medical University Hospital, Taipei Medical University, Taipei, Taiwan  
\*Corresponding author



@OHDSI

www.ohdsi.org

#JoinTheJourney



ohdsi



# #OHDSISocialShowcase This Week

## Wednesday

## Predicting outcome in emergency room patients with Suspected Gastrointestinal Infection using OMOP-CDM

(So Hee Lee, Byungjin Choi, Min Ho An, Junhyuk Chang, Harrin Kim, Rae Woong Park)



## Predicting outcome in emergency room patients with Suspected Gastrointestinal Infection using OMOP-CDM

So Hee Lee<sup>1</sup>, Byungjin Choi, MD<sup>2</sup>, Min Ho An, MD<sup>2</sup>, Junhyuk Chang<sup>1</sup>, Harrin Kim<sup>1</sup>, Sujin Gan, RN<sup>1</sup>, Rae Woong Park, MD, Ph.D<sup>1,2</sup>

<sup>1</sup>Department of Biomedical Sciences, Ajou University Graduate School of Medicine, Suwon, Republic of Korea

<sup>2</sup>Department of Biomedical Informatics, Ajou University School of Medicine, Suwon, Republic of Korea



### Background & Objectives

- Gastroenteritis is among the most frequently diagnosed conditions in emergency departments across the country
- Symptoms of gastroenteritis often begin mildly, with fever, diarrhea, abdominal pain, and vomiting. However, without prompt treatment, these symptoms can worsen, leading to more severe conditions that affect individuals of all ages
- In this study, we aim to predict 7-day 1) ER revisits, 2) ICU admissions, and 3) mortality, to assess the severity of gastroenteritis
- By identifying patients in high risk for disease prognosis, it enables quick triage and timely treatment, ultimately improving outcomes and reducing transmission

### Methods

#### Data sources

- Ajou University School of Medicine (AUSOM) database
  - Electronic health records (1994.01 ~ 2024.02)
  - OMOP-CDM v5.3.4

#### Study population

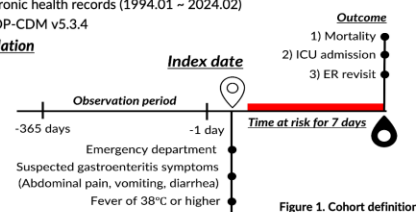


Figure 1. Cohort definition

#### Model development and evaluation

- Machine learning Algorithms:
  - Gradient boosting model (GBM) & Least absolute shrinkage and selection operator (LASSO)
- Covariates
  - Demographics, condition, drug, measurement and visit
  - Time frames : Long-term (-365 days) and Short term (-1 day) prior to the index date
  - The short-term period was chosen to capture recent conditions crucial for predicting the severity and treatment response of the patient
- Data split: Split into the train (75%) and test set (25%) in 3-fold cross validation
- Model performance
  - The area under the receiver operating curve (AUROC)
  - Youden index to determine threshold for high and low risk groups

#### Survival Analysis for ICU admission

- To assess the association between the risk of ICU admission and the incidence of 7-day mortality through the survival analyses
- Cox proportional model: for calculating the hazard ratio
- Kaplan-Meier method: for plotting the survival curve

### Results

- GBM outperformed LASSO in AUROC for predicting ER revisits, ICU admissions, and mortality (0.758 vs. 0.679, 0.964 vs. 0.947, and 0.990 vs. 0.980, respectively)

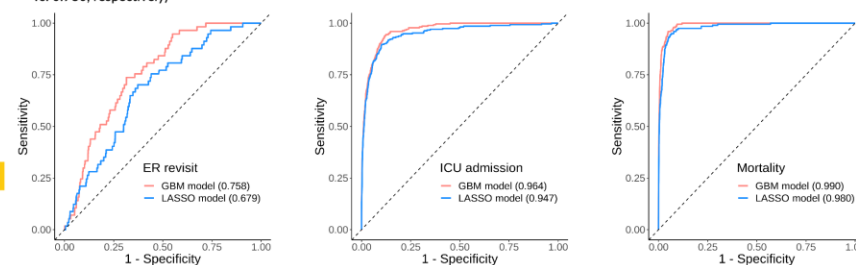


Figure 2. Model Performance for severity of gastroenteritis; (a) Prediction for ER revisit, (b) Prediction for ICU admission, (c) Prediction for mortality

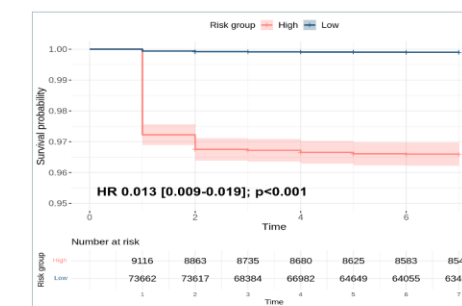


Figure 3. Kaplan-Meier survival analysis of Gradient boosting machine for ICU admission

- Based on the predicted results (Youden index: 0.005) by GBM prediction model in ICU admission, the low-risk group had a significantly lower hazard ratio for 7-day mortality (HR 0.013, 95% CI [0.009-0.019],  $p < 0.001$ )

### Conclusions

- We developed CDM-based prediction models to assess the severity of gastroenteritis outcomes, including ER revisits, ICU admissions, and mortality
- These models demonstrated moderate accuracy in predicting severity and distinguishing high-risk patients, aiding in timely treatment and improving patient outcomes

### Acknowledgements

This research was funded by a grant from the Korea Health Technology R&D Project through the Korea Health Industry Development Institute (KHIDI), funded by the Ministry of Health & Welfare, Republic of Korea (grant number: HR16C0001) and this research was supported by a Government-wide R&D Fund project for infectious disease research (GFID), Republic of Korea (grant number: HG22C0024, KH124685)



# #OHDSISocialShowcase This Week

## Thursday

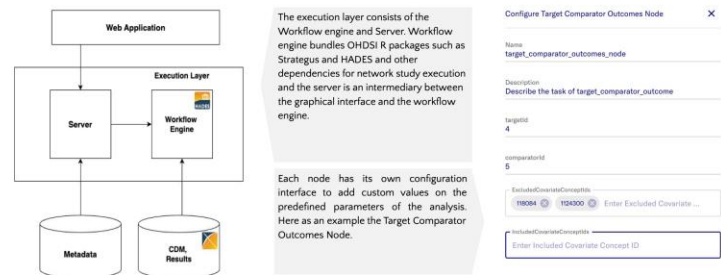
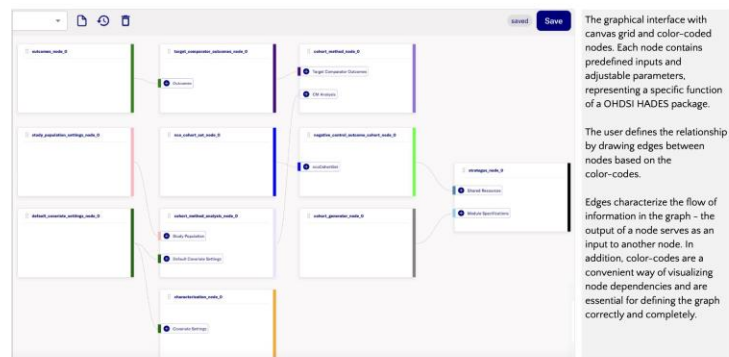
# A Graphical Interface and Workflow Engine for OHDSI Network Study design and execution

(Sivanaga Sai Krishna Santan Maddi, Hengxian Jiang, Peter Hoffmann)

Interactive, drag-drop interface with pre-installed and centralized execution layer for OHDSI network study design and execution

### Graphical Interface and Workflow Engine for Strategus

**Background:** The design and execution of network studies using R packages - Strategus and HADES - require prior knowledge in R programming and technical expertise. This limits participation to technical users only and becomes a barrier for non-technical users who are essential for collaborative research. We have developed a software that includes a user-friendly interface with an interactive drag-and-drop system complementing R packages, and an execution layer with an environment for executing network studies.



Sivanaga Sai Krishna Santan Maddi, Hengxian Jiang, Peter Hoffmann  
D4L data4life Asia Limited







# #OHDSISocialShowcase This Week

Friday

Explore the opinions and attitudes of the application of common data models in regional databases from the perspective of Chinese people

(Yexian Yu, Meng Zhang, Yongqi Zheng, Feng Sun)



## Explore the opinions and attitudes of the application of CDM in regional databases from the perspective of Chinese population

Yexian Yu<sup>1,2,3</sup>, Meng Zhang<sup>1,4</sup>, Yongqi Zheng<sup>1,4</sup>, Feng Sun<sup>2,3,4</sup>

<sup>1</sup> Hainan University, Haikou, China

<sup>2</sup> Hainan Boao Lecheng International Medical Tourism Pilot Zone Administration, Hainan Lecheng Institute of Real World Research, Lecheng, China

<sup>3</sup> Department of Epidemiology and Biostatistics, School of Public Health, Peking University, Beijing, China

<sup>4</sup> Key Laboratory of Epidemiology of Major Diseases (Peking University), Ministry of Education, Beijing, China

\* Corresponding Author: Dr. Feng Sun (sunfeng@bjmu.edu.cn)



### The role of CDM in the interconnection of healthcare data

CDM has become an effective tool for inter-regional and inter-institutional connectivity, integration, and collaborative analysis and utilization of health medical data, which plays a significant role in ensuring data consistency in multi-center cohort, epidemiological analysis, public health decision-making, and other research areas.

- The application of CDM in China is relatively low, which may be related to a lack of understanding and recognition of CDM<sup>1</sup>. Promoting and applying CDM in China requires more education and training to enhance the awareness of medical and health professionals about the importance of CDM.

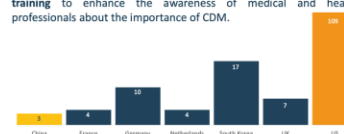


Figure 1. Distribution of Published Articles by Countries.

### Methods

- A questionnaire study was conducted to measure the acceptance of transitioning from regional databases to standardized CDMs among the Chinese population.
- The study explored participants' understanding of CDM and the Observational Medical Outcomes Partnership (OMOP), as well as their views on the necessity of CDM for regional databases in China.

Table 1. Questionnaire Items.

Questionnaire Items	
Gender	CDM Awareness
Age	OMOP Awareness Source
Working/studying Area	CDM Awareness Source
Occupation	OMOP Mainstream Ability
Education Level	CDM Necessity
	OMOP Benefits
	CDM Type
	OMOP Challenges
	OMOP Awareness
	Suggestions

- Analysis of the survey results aims to uncover the current state, challenges, and trends of CDM implementation in the Chinese medical field, offering a basis for future data standardization and sharing.



\* All statistical analyses were performed using R version 4.4.1.

Yexian Yu: yuyexian@hainanu.edu.cn

Contact: contact@ohdsi.org

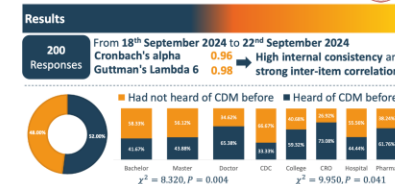


Figure 2. Proportion of CDM Awareness in different factors.

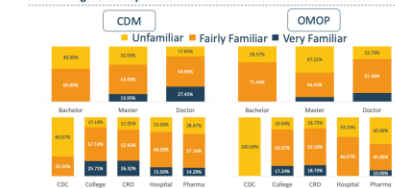
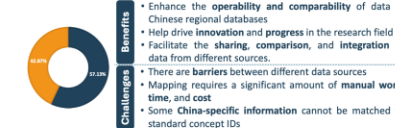


Figure 3. Association between Familiarity and Different Factors.

- More than half of the participants believe that OMOP could become the mainstream choice for CDM in Chinese regional databases.



### Conclusions

- Based on the statistical analysis of the questionnaire results, participants generally hold a positive attitude towards the application of CDM in regional databases in China.
- They also offer their suggestions, which provide valuable guidance and direction for the future promotion of CDM applications in the healthcare sector in China.



### References

<sup>1</sup> Reinecke L, Zoch M, Reich C, Sedlmayr M, Bathelt F. The Usage of OHDSI OMOP - A Scoping Review. Stud Health Technol Inform. 2021 Sep 21;283:95-103.

### Acknowledgment

The authors would like to thank the sponsorship from the Joint Real-World Evidence Research Lab founded by Peking University Health Science Center-AstraZeneca established since Oct.2022, this research is part of the internship program starting July.2024.

<https://news.bion.com/article/cecb829e26a8.html>.



@OHDSI

[www.ohdsi.org](http://www.ohdsi.org)

#JoinTheJourney



ohdsi





# Where Are We Going?

**Any other announcements  
of upcoming work, events,  
deadlines, etc?**



# Three Stages of The Journey

**Where Have We Been?**

**Where Are We Now?**

**Where Are We Going?**





# June 17: ATLAS Deepdive

## Cohorts and Concept Sets



## Christopher Knoll

Director, Observational Health Data Analytics  
Janssen Research and Development



## Richard Boyce

Associate Professor, Department of Biomedical  
Informatics  
University of Pittsburgh

Join us  
throughout June  
to help create the  
roadmap for  
ATLAS!



# Week 3 ATLAS Survey:

## Cohorts and Concept Sets



**These weekly surveys  
will help us build future  
versions of ATLAS!**

**We are asking for input  
throughout our global  
community!**





**The weekly OHDSI community call is held  
every Tuesday at 11 am ET.**

**Everybody is invited!**

**Links are sent out weekly and available at:  
[ohdsi.org/community-calls-2025](https://ohdsi.org/community-calls-2025)**