

# OHDSI2025 Global **Symposium Preview**

**OHDSI Community Call** Sept. 9, 2025 • 11 am ET









# **Upcoming Community Calls**

Date	Topic				
Sept. 9	Global Symposium Preview				
Sept. 16	OHDSI/OMOP Research Spotlight				
Sept. 23	Educating on OHDSI: Lessons Learned				
Sept. 30	OHDSI 2025 Poster Preview Mad Minutes / Symposium Logistics				
Oct. 7	No Call – OHDSI Symposium				
Oct. 14	Welcome to OHDSI				
Oct. 21	Meet the Titans				









# Sept 16: OHDSI/OMOP Research Spotlight



**Jessie Tong** 

Assistant Professor, Johns Hopkins University

Unlocking efficiency in real-world collaborative studies: a multi-site international study with one-shot lossless GLMM algorithm • NPJ Digital Medicine



### Kim López Güell

Dphil Candidate, University of Oxford

Clusters of post-acute COVID-19 symptoms: a latent class analysis across 9 databases and 7 countries • Journal of Clinical Epidemiology



### Jen Wooyeon Park

PhD Student, Johns Hopkins University

Breaking data silos: incorporating the DICOM imaging standard into the OMOP CDM to enable multimodal research • JAMIA



### **Abigail Newbury**

PhD Student, Columbia University

Multi-domain rule-based phenotyping algorithms enable improved GWAS signal • NPJ Digital Medicine



### **Benjamin Martin**

Postdoctoral Fellow, Johns Hopkins University

Identification of Adult Dermatomyositis Patients Using Real-World Data Sources • Arthritis Care and Research









# Three Stages of The Journey

Where Have We Been? Where Are We Now? Where Are We Going?









## **OHDSI Shoutouts!**



Congratulations to the team of Junqing Xie, Mike Du, Yuchen Guo, Cesar Barboza, James T Brash, Antonella Delmestri, Talita Duarte-Salles, Jasmine Gratton, Romain Griffier, Raivo Kolde, Wai Yi Man, Núria Mercadé-Besora, Marek Oja, Sarah Seager, Katia Verhamme, Dina Vojinovic, Edward Burn, Daniel Prieto-Alhambra, Martí Català, Annika M Jödicke on the publication of Trends in prescription opioid use in Europe: A DARWIN EU® multinational cohort study including seven European countries in Frontiers in Pharmacology.



TYPE Original Research PUBLISHED 18 August 2025 DOI 10.3389/fphar.2025.1608051



#### **OPEN ACCESS**

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Xie J. Du M. Guo Y. Barboza C. Brash JT. Delmestri A, Duarte-Salles T, Gratton J, Griffier R, Kolde R, Man WY, Mercadé-Besora N Oja M, Seager S, Verhamme K, Vojinovic D, Burn E, Prieto-Alhambra D, Català M and Jödicke AM (2025) Trends in prescription opioid use in Europe: A DARWIN EU® multinational cohort study including seven European countries. Front. Pharmacol. 16:1608051 doi: 10.3389/fphar.2025.1608051

© 2025 Xie, Du, Guo, Barboza, Brash, Delmestri Duarte-Salles, Gratton, Griffier, Kolde, Man, Mercadé-Besora, Oia, Seager, Verhamme Vojinovic, Burn, Prieto-Alhambra, Català and

Trends in prescription opioid use in Europe: A DARWIN EU® multinational cohort study including seven European countries

Junging Xie<sup>1†</sup>, Mike Du<sup>1†</sup>, Yuchen Guo<sup>1</sup>, Cesar Barboza<sup>2</sup>, James T. Brash<sup>3</sup>, Antonella Delmestri<sup>1</sup>, Talita Duarte-Salles<sup>2,4</sup>, Jasmine Gratton<sup>3</sup>, Romain Griffier<sup>5</sup>, Raivo Kolde<sup>6</sup>, Wai Yi Man<sup>1</sup>, Núria Mercadé-Besora<sup>4</sup>, Marek Oja<sup>6</sup>, Sarah Seager<sup>3</sup>, Katia Verhamme<sup>2</sup>, Dina Vojinovic<sup>7</sup>, Edward Burn<sup>1</sup>, Daniel Prieto-Alhambra 1.2\*, Martí Català 1 and Annika M. Jödicke 1

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Background: The opioid crisis has been a serious public health challenge in North America for decades, despite numerous efforts to mitigate its devastating consequences. As concerns grow about a similar situation developing in Europe, we evaluated the trends in opioid use and characterized prescribing indications across seven European countries.











## **OHDSI Shoutouts!**



German Medical Data Sciences 2025: GMDS Illuminates Health R. Röhrig et al. (Eds.) © 2025 The Authors. This article is published online with Open Access by IOS Press and distributed under the terms of the Creative Commons Attribution Non-Commercial License 4.0 (CC BY-NC 4.0)

Development and Implementation of an Open, Modular, and Participatory Toolchain for Distributed IT Development in Healthcare Research - Lessons Learned

Daniel NEUMANN<sup>a,1</sup>, Richard GEBLER<sup>b</sup>, Jana KIEDERLE<sup>c</sup>, Jördis BECK<sup>d</sup>, Fabio AUBELE<sup>c</sup>, Alexander STRUEBING<sup>a</sup>, Florian SCHMIDT<sup>a</sup>, Matthias REUSCHEa, Helene KOESTERf, Markus LOEFFLERa, and Sebastian STAEUBERT

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Abstract. Introduction: Distributed healthcare research infrastructures face significant challenges when translating routine clinical data into harmonized,

research-ready formats using HL7 FHIR standards. State of the Art: Existing FHIR-based pipelines such as the SMART/HL7 FHIR Bulk Data Access API, FHIR-to-OMOP mappings, and analytical services like Pathling demonstrate technical feasibility. However, most assume semantically valid FHIR data, operate within single-institution settings, and lack practical guidance for deployment across heterogeneous, regulated environments. Technical Framework and Deployment: Within the German Medical Informatics Initiative (MII) and the INTERPOLAR project, we developed an open, modular, and participatory toolchain for decentralized FHIR-based data transformation and export across multiple Data Integration Centers (DICs). The toolchain supports FHIR extraction, profile-based transformation, REDCap integration, and OMOP-compatible export. Deployment required adapting to local infrastructures, regulatory boundaries (e.g., de-identified FHIR stores, restricted network access), and clinical domain needs. Configurable modules, proxy support, and site-specific adaptations were essential for integration into operational hospital workflows. Lessons Learned: Key lessons include the necessity of early access to real data, the limitations of synthetic test data, the value of joint workshops for profile interpretation, and the need for adaptable validation tooling. Organizational knowledge gaps, inconsistent FHIR implementations, and performance issues in resource flattening were addressed through co-design and

Congratulations to the team of Daniel Neumann, Richard Gebler, Jana Kiederle, Jördis Beck, Fabio Aubele, Alexander Struebing, Florian Schmidt, Matthias Reusche, Helene Koester, Markus Loeffler, Sebastian Staeubert on the publication of Development and Implementation of an Open, Modular, and Participatory Toolchain for **Distributed IT Development in Healthcare** Research – Lessons Learned in Volume 331 of Studies in Health Technology and Informatics: German Medical Data Sciences 2025: GMDS Illuminates Health.









# 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/WebAPI			
Tuesday	12 pm	Generative AI and Analytics			
Tuesday	3 pm	Oncology Outreach/Research Subgroup			
Wednesday	9 am	Patient-Level Prediction			
Wednesday	2 pm	Natural Language Processing			
Wednesday	7 pm	Eyecare and Vision Research			
Thursday	7 am	Europe Community Call			
Thursday	9:30 am	Network Data Quality			
Thursday	10 am	Rare Diseases			
Thursday	10:30 am	Evidence Network			
Friday	9 am	Phenotype Development & Evaluation			
Friday	10 am	GIS – Geographic Information System			
Friday	11 am	Clinical Trials			
Friday	11:30 am	Steering			
Friday	11 pm	China Chapter			
Monday	9 am	Vaccine Vocabulary			
Monday	10 am	Africa Chapter			
Monday	10 am	Getting Started Subgroup			
Monday	11 am	Data Bricks User Group			
Monday	2 pm	Electronic Animal Health Records			
Tuesday	10 am	CDM Survey Subgroup			









# **2025 Europe Community Calls**

Date	Topic
Sept. 11	Europe Community Call Introduction / DARWIN EU Update
Oct. 9	TBA
Nov. 13	Patient-Reported Outcome Measures (PROMs)
Dec. 11	Vocabularies in Europe















## **Science Summit 2025**

alongside the United Nations General Assembly (UNGA80)

9 – 26 September 2025



https://sciencesummitnyc.org/

### Science for a Sustainable Future: **Showcasing Science** Collaboration

The role and contribution of science in attaining the United Nations Sustainable Development Goals (SDGs) will be the central theme of the Science Summit. The objective is to enable science collaborations to demonstrate how science supports the attainment of the UN SDGs and Agenda 2030.

The Summit will examine what enabling policy, regulatory and financial environments are needed to implement and sustain the science mechanisms required to support genuinely global scientific collaborations across continents, nations and themes.

Scientific discovery through the analysis of massive data sets is at hand. This data-enabled approach to science, research and development will be necessary if the SDGs are to be achieved.

SCIENCE FOR GLOBAL CHALLENGES  $\rightarrow$ 

Full programme is <u>here</u>













### Standardizing Health Data and Analytics to Accelerate Clinical Impact and Global Reach: Part 1

◆ Theme: Digital / All



Observational Health Data Science and Informatics (OHDSI) is a global community that uses data harmonized to the OMOP Common Data Model, standardized vocabulary, data quality checks and validated analytics to produce rigorous evaluation of big data from existing health databases. Through sharing of computer codes and summary statistics instead of patient-level data, OHDSI preserves privacy while enabling collaboration across institutions, countries, and continents. Large-scale, real-world studies through OHDSI network collaborations have revealed valuable insights into clinical care and public health.

#### Speakers:



Agnes Kiragga Global Health Leader,...

Organization: African Population Health and Research Centre



Chan Seng You Assistant Professor

Organization: Yonsei University College of Medicine



NicolePratt Professor, Biostatistic...

Organization: University of South Australia



George Hripcsak Professor, Biomedical...

Organization: Columbia University

Register

Session details











### Standardizing Health Data and Analytics to Accelerate Clinical Impact and Global Reach: Part 2

◆ Theme: Digital / Al



Observational Health Data Science and Informatics (OHDSI) is a global community that uses data harmonized to the OMOP Common Data Model, standardized vocabulary, data quality checks and validated analytics to produce large-scale evaluation of real world data. Through sharing of computer codes and summary statistics instead of patient-level data, OHDSI preserves privacy while enabling collaboration across institutions, countries, and continents. Large-scale, real-world studies by OHDSI members have revealed valuable insights into clinical care and public health.

#### Speakers:



Cynthia Sung Adjunct Associate...

Organization: Duke-NUS Medical School Centre of Regulatory Excellence



Patrick Ryan /P Janssen...

Organization: OHDSI Observational Health Data Science and Informatics



Katia Verhamme Associate Professor of...

Organization: Erasmus University Medical Center



Peter Rijnbeek Professor, Medical...

Organization: Erasmus University Medical Center



Iulio Oliveira

Organization: Precision Data











### Registration links

Part 1 Sep 18, 8:30-10:30 EDT: <a href="https://event.sciencesummitnyc.org/list-of-sessions/detail/131">https://event.sciencesummitnyc.org/list-of-sessions/detail/131</a>

Part 2: Sep 18, 11:00-13:00 EDT <a href="https://event.sciencesummitnyc.org/list-of-sessions/detail/130">https://event.sciencesummitnyc.org/list-of-sessions/detail/130</a>

Full programme here: <a href="https://event.sciencesummitnyc.org/list-of-sessions">https://event.sciencesummitnyc.org/list-of-sessions</a>

## Part 1 (8:30 am ET)

- 1. Observational Health Data Science and Informatics (OHDSI): Inclusive and Collaborative Science. George Hripcsak
- 2. Promoting Data Harmonization and Data Science in Africa. Agnes Kiragga
- Rapid Response to the Covid-19 Pandemic Using a National Scale Database. Chan Seng You
- OHDSI in Asia and the Pacific Rim. Nicole Pratt
- 5. Q&A Session

## Part 2 (11 am ET)

- 1. Enabling Reliable Evidence Generation from Real-word Data in Europe. Peter Rijnbeek
- 2. DARWIN-EU® Delivering Real World
  Evidence to Support Regulatory Decisionmaking by the European Medicines Agency.
  Katia Verhamme
- 3. OHDSI Adoption and Current Implementation Landscape in Latin America. Julio Cesar Barbour Oliveira
- 4. Learning Opportunities for OHDSI Skills Development. Cynthia Sung
- Clinical and Public Health Impact of OHDSI. Patrick Ryan
- 6. Q&A Session











# **Titan Award Nominations Are Due TONIGHT!**

The Titan Awards have been handed out annually since 2018 to recognize OHDSI collaborators (or collaborating institutions) for their contributions towards OHDSI's mission.

Nominations for the 2025 Titan Awards are now open. Please complete your nominations by our Sept. 9 (8 pm ET) deadline!

ohdsi.org/titan-awards











# Upcoming Presentation (Sept. 29, 1-2 pm ET)

# Delivery, Not Hype: How to Harmonise FHIR × openEHR × OMOP in practice

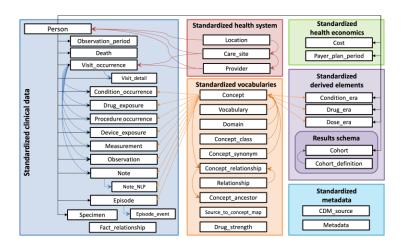
A Health Futures Collective Series - powered by Evidentli, Fuzzy, openEHR, Professional Development and Summer Programmes, Institute of Extended Learning at Imperial College

### **Confirmed panellists**

Grahame Grieve — Inventor of FHIR Rachel Dunscombe — CEO, openEHR Foundation Davera Gabriel — Lead Author, FHIR to OMOP IG Guy Tsafnat — Expert in real-world data & Al

### Moderator

Ram Rajaraman — Healthcare & Life Sciences Lead, Quantexa















# Jamie Weaver Scholarship at University of Oxford

**PhDs** ○ Improving the quality of real world evidence by measuring and minimising outcome misclassification using the OMOP common data model and large multinational health data (Botnar-2025-8) University of Oxford > Botnar Research Centre Prof Dani Prieto-Alhambra Tuesday, December 02, 2025 Funded PhD Project (Students Worldwide)

#### **About the Project**

This scholarship and work has been proposed to continue and expand work started by the late James (Jamie) Weaver. Jamie was a talented and bright data scientist and DPhil student working with us on the use of methods to minimise the impact of outcome misclassification in real world evidence (RWE). Funding has been secured, from the Medical Sciences Division, Brasenose College, and NDORMS, for this project to continue his important work on this extremely relevant topic; the successful candidate will be assigned to Brasenose College.

Real world evidence (RWE) is generated by leveraging and processing large routinely collected health data. Despite difficulties in the analysis of such information for causal inference purposes, RWE has recently been shown as a reliable source of data when used using adequate methods for trial emulation [1, 2]. We participate in multiple European and international networks to generate reliable information to inform, amongst others, regulatory decision making and health technology assessments.

Through ongoing collaborations, we leverage multiple international datasets mapped to the Observational Medical Outcomes Partnership (OMOP) Common Data Model in a federated manner. Previous work led by our student Jamie Weaver uncovered the impact of outcome misclassification on the estimation of background rates of adverse events, and proposed new methods to account for this in future studies [3].

Through this 3-year PhD funded studentship, we aim to investigate how novel methods can be applied to measure and account for outcome misclassification in RWE studies. by researching:

- 1. The use and application of artificial intelligence (and specifically large language models) for the generation and validation of computable phenotypes
- 2. The impact of outcome misclassification in different data assets
- 3. The performance of existing and novel methods to account for outcome misclassification in international RWE studies











# Global Symposium: Oct. 7-9



ohdsi.org/ohdsi2025



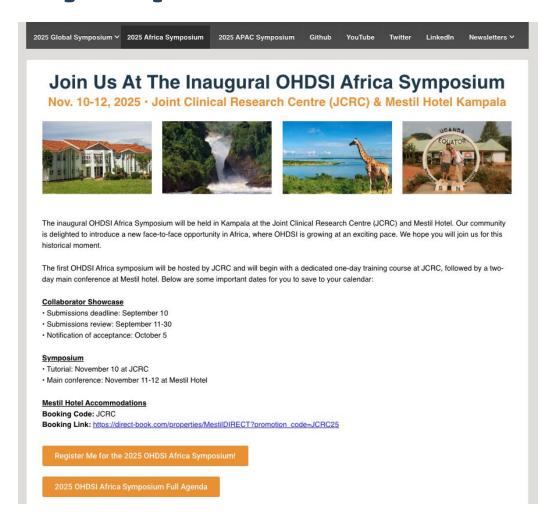








# Africa Symposium: Nov. 10-12



ohdsi.org/africa2025











# APAC Symposium: Dec. 6-7

The 2025 OHDSI APAC Symposium will be held Dec. 6-7 in Shanghai, China at the Shanghai Jiao Tong University. It will feature a 1-day tutorial and a 1-day main conference. Here are some important dates for you to save to your calendar:



### **Collaborator Showcase**

- Submissions deadline: September 7
- •Submissions review: September 8 October 9
- Notification of acceptance: October 17



ohdsi.org/apac2025











# Monday

ECRAID-Base OMOP-CDM ETL Architecture for Scalable Health Data Integration

(Panagiotis Gialernios, Shirah Cashriel, Marc Padros Goossens, Frank Leus, Freija Descamps, Lauren Maxwell, Ankur Krishnan)

### A Modular ETL Framework for Transforming Multi-Source Medical Data to OMOP Common Data Model

Title: Architecture for Scalable Health Data Integration

Background: ECRAID-Base is an EU-funded project aiming to efficiently generate rigorous evidence to improve the diagnosis, prevention, and treatment of infections, while responding to emerging infectious diseases and antimicrobial resistance threats. At its core are 5 perpetual observational studies (POS), with POS-VAP (Ventilator Associated Pneumonia) harmonized to MOP-CDM through the EHDEN (European Health Data and Evidence Network) project. To optimize the harmonization process, common medical reference data such as pathogen and virus lists are mapped once as shared components and reused across all POS studies ensuring consistency while reducing redundant mapping efforts.

Methods: The ECRAID-Base data is collected in a question/answer format, with each pair mapped to OMOP-CDM using expert-defined instructions that specify target tables, fields, and concept IDs.

#### Modular ETL Architecture Overview



Modular approach enables easy addition of new data sources while maintaining consistency

Each data source maintains its specific mappings while leveraging common instructions

#### Key Updates to ECRAID-Base ETL Codebase

Codebase Restructuring

- Modular Imput

- Chair separation of source-specific logic
- Improved sosiability

- Reduced human error

- Reduced human error

- Reduced human error

- Reduced human error

Framework Benefits

ensure source-specific changes don't impact other day

/ Isolated components ensure source-specific changes don't impact other data source

√ New data sources are easier to add

√ Streamlined integration while maintaining scalability

Conclusion: Our modular ETL architecture significantly enhances scalability, maintainability, and adaptability for integrating heterogeneous medical data into OMOP CDM. The framework's success with harmonized eCRFs (electronic Case Report Forms) under ECRAID-Base demonstrates the importance of broader harmonization acr medical data sources, streamlining management and ensuring adaptability to evolving health data challenges.





Panagiotis Gialernios¹, Shirah Cashriel¹, Marc Padros Goossens² Frank Leus², Freija Descamps¹, Lauren Maxwell², Ankur Krishnan delangen Allians Goossen Allians an Infertiona Disease (ECRAID)















## Tuesday

Multiple myeloma lines of treatment: from drug\_exposure to proper Episode table

(Dmytro Dymshyts, Rupa Makadia, Laura L. Hester)

### Multiple myeloma lines of treatment: from drug\_era to proper OMOP **Episode table**

How to extract Multiple Myeloma (MM) treatment regimen from drug\_era into Episode table

Background: The drug data can be abstracted on the 4 principal levels: drug exposure (single drug administration or prescription), drug era (continuous drug administration or prescription), treatment regimen (drugs used in combination with fixed schedule), line of therapy (several regimens used consecutively united by one clinical intent). The first two levels exist in our OMOP common data model (CDM) datasets, and we need an effective way of capturing and storing lines and regimen.

#### Main principles applied: Figure 1. Abstraction of drug era into Regimen and line of therapy in individual patient 1. First regimen starts as any MM-specific drug exposure after MM diagnosis 2. All MM-specific drugs within 30 days of regimen start considered a part of regimen (applicable to first and subsequent regimen) 3. Regimen ends if either a.New drug is added b.Drug is removed (not used for more than 90 days 4. New regimen starts as either a. Next day of the previous regimen end b.Start of the new drug era, if there was gap between previous regimen 5. Maintenance therapy is a monotherapy duration >60 days that follows a combination therapy that lasts more than 30 days 6. Regimens are grouped into line of therapy when they

Table 1.Characterization of treatment regime

	Flatiron MM	Optum EHR	Optum SES	HealthVerity
total number of patients	17433	39311	44696	71005
median count of regimen	2	2	2	2
median count of lines	1	1	- 1	2
median length of regimen, days	66	56	89	85
median length of line, days	126	85	129	132
most frequent line of therapy	bortezomib, lenalidomide, dexamethasone	lenalidomide, dexamethasone (with bortezomib, lenalidomide, dexamethasone as the next most common)	bortezomib, lenalidomide, dexanethasone	bortezomib, lenalidomide, dexamethasone

Regimen and lines are consistent across the databases

Conclusion: The algorithm creates two additional levels of abstraction of multiple myeloma treatment: Regimen and Lines of therapy, - which are useful in oncological observational research. It shows consistent and plausible results in US databases, with our next steps to abstract such regimen for our in-house European databases as well as sharing the approach with OHDSI data partners across Europe

a.Stem cell transplant and related

previous regimen start

treatment regimen

(conditioning/lymphodepletion) therapy b.Apheresis, anti-plasma cell treatment, CAR-T

c.Regime and its corresponding maintenance d.Addition of immunomodulatory or proteasome inhibitor drugs within 90 days of the

7. The regimen are mapped to HemOnc concepts by matching ingredients and populate EPISODE episode\_object\_concept\_id

8. Custom concepts were created to support the line of

9. The line of treatment becomes a parent episode of



Dmytro Dymshyts, Rupa Makadia, Laura L. Hester

Johnson &Johnson













# Wednesday

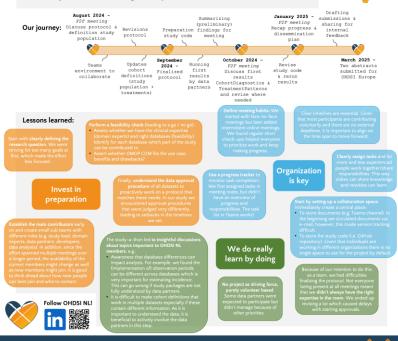
Lessons learned from the OHDSI-NL study-a-thon on breast cancer

(Aniek F. Markus, Sofia Bazakou, Renske Los, Julia Kurps, Jelle Evers on behalf of OHDSI NL) The OHDSI NL study-a-thon has been a fun and productive way to learn together, but we need to expand our network of data partners in the Netherlands!

Lessons learned from the OHDSI NL study-a-thon on breast cancer

Background: To foster collaborations on both national and global level, OHDSI national nodes have been set up in several countries. The Dutch node (OHDSI NL) initiated a network study to assess the extent to which we are ready to execute network studies and answer clinical questions at a national level in the Netherlands. In this work, we reflect on our findings and share the key lessons learned throughout the process.





Sofia Bazakou, Julia Kurps, Jelle Evers, Renske Los, Aniek Markus on behalf of OHDSI NL













# **Thursday**

Catalysing biobank diversity analysis through a common data model

(Karyn Mégy, Ben Hollis, Celia Burgos, Katherine R Smith, Prasad Gunasekaran, Sean O'Dell, Sebastian Wasilewski, Quanli Wang, Slavé Petrovski)

### Catalysing biobank diversity analysis through a common data model.

Karyn Mégy¹, Ben Hollis¹, Celia Burgos Sequeros¹, Katherine R. Smith¹, Prasad Gunasekaran¹, Sean O'Dell¹, Sebastian Wasilewski¹, Quanli Wang², Slavé Petrovski¹.

1. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics Research, Discovery Sciences, BioPharmaceuticals R&D, AstraZeneca, Cambridge, UK. 2. Centre for Genomics R&D, AstraZeneca, Centre for Genomics R&D, AstraZe

AstraZeneca's Centre for Genomics Research (CGR) is building one of the world's most comprehensive and entically diverse genetic resources by integrating genetic and phenotypic data from multiple biobanks. We are expanding the ancestral diversity of our datasens souring the data used in our research reflects global populations, so our science is designed and able to benefit a broad range of people. However, to fully leverage the power of this ethnic diversity, it is essential to standardise the data into a common data model.

UK Biobank (UKB) is one of the golden standard, in terms of data diversity, sources but also coding systems. However, it is very reflective of the UK population and health care system. Biobanks from different countries use multiple clinical coding systems, different units (e.g. Hba1c: mmol/mol vs. %), medication names (e.g. metformin vs. metformina), and local language. This makes their comparison very challenging without any standardisations of format and content.

den	Iltiple standards  Source of health data available in our largest cohorts						
ery alth ent ling alc: e.g. ocal ery s of		Hospital data	Primary care	Cancer data	Questionnaires	Free text	And also
	UK Biobank	WHO ICD9 & 10	Read2 & 3	yes	formatted		Lab. tests, procedures, multi-omics, drugs
	US cohort #1	CM ICD9 & 10, SNOMED	CM ICD9 & 10, SNOMED	CM ICD9 & 10	-	-	Lab. tests, procedures, drugs
	Mexico cohort #1	-	-	WHO ICD10	WHO ICD10, formatted	yes	Lab. Tests, drugs (in Spanish), Metabolomic
	Pakistan cohort #1	-	-		Ad hoc data dictionary	yes	-
	ICD: International Classific	ation of Diseases, versio	ns 8, 9 and 33, in the WHO	or US (CM) system			

#### Tailored vs. large biobanks

Smaller tailored biobanks are focused on specific diseases or populations.

#### Clinical data

- · Questionnaires, not derived from electronic health records
- Collected at the point of recruitment
- Missingness and error-prone (typos)
- · => non-standard OMOP concepts, or no concepts

Format of the data, or even of the questions, can change from one release to the next (e.g. 'Gender: male / female' vs. 'Gender male: yes / no).

Unique phenotypic data not typically gathered across other biobank (e.g. types of smoking/chewing tobacco) prompting considerations on the extent of original data that should be converted to OMOP.

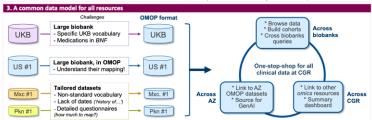
Large biobanks are disease-agnostic, with a broad spectrum of diagnoses, medications, and measurements.

#### Clinical data

- Sourced from electronic health records and optionally additional targeted questionnaires
- Well-structured, minimal missing values, systematic and automatic collection

Format remains largely consistent between releases.

However, harmonising medications or measurements poses challenges due to inconsistencies in data collection, such as varying units within the same measurement concept, or the lack of an international standard for drugs (e.g. Intent in British National Formulary –BNF- vs. active substance in Anatomical Therapeutic Chemical classifications –ATC-)



#### 4. Take home messages

- > Harmonising data format and content is essential for maximising the value of multiple biobanks.
- Effective transformation strategies depend on a deep understanding of each dataset
- > Lack of standards in the scientific community (e.g. measurements, medications) can limit the mapping.
- > Taking a holistic view and considering the broader context is key when standardising multiple datasets.

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AstraZeneca 2





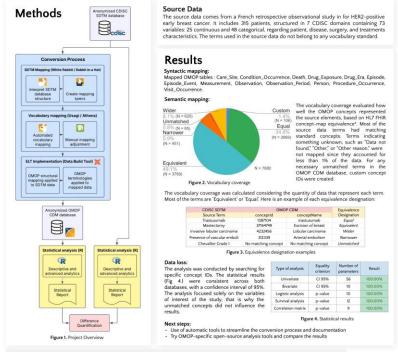
# **Friday**

Key learnings after transformation of a CDISC SDTM database into OMOP CDM

(Amélie Lambert, Claire Castagne, Jacek Chmiel, Eric Boernert, François Margraff, David Pau, Camille Bachot, Lukasz Kaczmarek, Dimitar Toshev, Thomas Stone)

## Key learnings after transformation of a CDISC SDTM database into OMOP CDM

Background: Interoperability between databases is becoming increasingly important to facilitate analyses from multiple sources. The OMOP Common Data Model (CDM) is widely used to standardize the structure and content of databases, enabling large-scale observational studies. Converting existing databases to the OMOP format is essential for leveraging its benefits. The aim is to assess the statistical and scientific usefulness<sup>1,2</sup> of a transformation from CDISC Study Data Tabulation Model (SDTM) to OMOP CDM.



Key learnings: Our study showed that most of our database could be mapped to the OMOP CDM. The model is proficient at representing positive information such as what a patient has or experiences. However, representing negative information (what a patient does not have) requires additional effort and consideration, as the OMOP CDM is not designed for this purpose. Information loss during conversion varies based on the original database's level of detail and the mapping approach. Successful adaptation to the CDM necessitates modifications in both data generation and statistical analysis scripts. This involves ensuring that the terminology used in data collection matches the standardized vocabularies defined by OMOP, such as SNOMED, LOINC, and RxNorm. By doing so, it facilitates smoother data integration, reduces information loss, and enhances the accuracy of subsequent analysis.



Pau D, Bacher C, Montell C, Vine L, Bescher M, Sella M, (go) R. Comparison of anonymisation schwiques regarding seasing productibility. PLOS Digital Health. 2025 Feb 34(5):e0000735; 1/9gau R. Bachet C, Montell C, Boorber E. Christil I, Boocher M, Reu D, Cambridge L, Sella M, Control C, Sella M, Control C,



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# 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?









# 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





