

Connections for Future Collaborations

OHDSI Community Call Jan. 14, 2025 • 11 am ET

#JoinTheJourney





Upcoming Community Calls

Date	Topic
Jan. 14	Connnections for Future Collaborations
Jan. 21	Clinical Guideline Review, Session I
Jan. 28	Clinical Guideline Review, Session II
Feb. 4	First Week of 2025 Workgroup OKRs/Phenotype Phebruary







Three Stages of The Journey

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









Congratulations to the team of Melissa Finster, Maxim **Moinat and Elham** Taghizadeh on the publication of ETL: From the German **Health Data Lab data formats** to the OMOP Common Data Model in PLOS One.

PLOS ONE

RESEARCH ARTICLE

ETL: From the German Health Data Lab data formats to the OMOP Common Data Model

Melissa Finster 61*, Maxim Moinat2, Elham Taghizadeh 61

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- * melissa.finster@mevis.fraunhofer.de

Abstract

Objective

The German Health Data Lab is going to provide access to German statutory health insurance claims data ranging from 2009 to the present for research purposes. Due to evolving data formats within the German Health Data Lab, there is a need to standardize this data into a Common Data Model to facilitate collaborative health research and minimize the need for researchers to adapt to multiple data formats. For this purpose we selected transforming the data to the Observational Medical Outcomes Partnership Common Data Model.



A OPEN ACCESS

Citation: Finster M, Moinat M, Taghizadeh E (2025) ETL: From the German Health Data Lab data formats to the OMOP Common Data Model. PLoS ONE 20(1): e0311511. https://doi.org/10.1371/ journal.pone.0311511





Congratulations to the team of Martijn Schuemie, Anna Ostropolets, Aleh Zhuk, Uladzislau Korsik, Seung In Seo, Marc Suchard, **George Hripcsak and Patrick Ryan** on the publication of **Standardized** patient profile review using large language models for case adjudication in observational research in NPJ Digital Medicine.

npj | digital medicine

Article

Published in partnership with Seoul National University Bundang Hospital



https://doi.org/10.1038/s41746-025-01433-4

Standardized patient profile review using large language models for case adjudication in observational research

Check for updates

Martijn J. Schuemie ® ¹.².3 ⋈, Anna Ostropolets¹.⁴, Aleh Zhuk¹.⁵, Uladzislau Korsik¹.⁵, Seung In Seo¹.⁶, Marc A. Suchard¹.³, George Hripcsak¹.⁴ & Patrick B. Ryan¹.².⁴

Using administrative claims and electronic health records for observational studies is common but challenging due to data limitations. Researchers rely on phenotype algorithms, requiring labor-intensive chart reviews for validation. This study investigates whether case adjudication using the previously introduced Knowledge-Enhanced Electronic Profile Review (KEEPER) system with large language models (LLMs) is feasible and could serve as a viable alternative to manual chart review. The task involves adjudicating cases identified by a phenotype algorithm, with KEEPER extracting predefined findings such as symptoms, comorbidities, and treatments from structured data. LLMs then evaluate KEEPER outputs to determine whether a patient truly qualifies as a case. We tested four LLMs including GPT-4, hosted locally to ensure privacy. Using zero-shot prompting and iterative prompt optimization, we found LLM performance, across ten diseases, varied by prompt and model, with sensitivities from 78 to 98% and specificities from 48 to 98%, indicating promise for automating phenotype evaluation.







Congratulations to the team of Noah Hong, Yeh-Hee Ko, Jeong Hyun Park, Eun Jin Ha, Sung Ho Lee, Kang Min Kim, Hyun-Seung Kang, Jeong Eun Kim, Kwangsoo Kim, and Won-Sang Cho on the publication of Impact of **Uric Acid Levels on Mortality and** Cardiovascular Outcomes in Relation to Kidney Function in the Journal of Clinical Neuroscience.

Journal of Clinical Neuroscience 133 (2025) 111039



Contents lists available at ScienceDirect Journal of Clinical Neuroscience



journal homepage: www.journals.elsevier.com/journal-of-clinical-neuroscience



A common data model for oral anticoagulants-related risk of spontaneous intracranial hemorrhage

Noah Hong a,1 , Yeh-Hee Ko b,1 , Jeong Hyun Park c,d , Eun Jin Ha e,g , Sung Ho Lee g , Kang Min Kim g , Hyun-Seung Kang g , Jeong Eun Kim g , Kwangsoo Kim f,* , Won-Sang Cho g,f

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ARTICLE INFO

Neworth. Warfarin Non-vitamin K antagonist oral anticoagulants Risk factors

Spontaneous intracranial hemorrhage Common data model

ABSTRACT

Spontaneous intracranial hemorrhage (sICH) is a major complication associated with oral anticoagulation which results in a high mortality rate, and the incidence of anticoagulant-induced sICH has increased markedly, so it is necessary to investigate the risk of anticoagulanton-related sICH in a real-world setting. We aimed to investigate the incidence and risk factors of oral anticoagulant-related sICH using a common data model (CDM), and to determine whether a clinical study using the CDM would be comparable to conventional studies. After converting the various clinical codes of 12,821 patients taking oral anticoagulants, such as warfarin and non-vitamin K antagonist oral anticoagulants (NoACs), into the Observational Medical Outcomes Partnership (OMOP) CDM format, we analyzed the incidence and risk factors of sICH. sICH occurred in 0.5 % of 5,626 patients with warfarin and 0.2 % of 7,195 patients with NoAC. The mean duration of warfarin and NoACs before sICH occurrence was 251.4 ± 373.6 and 124.2 ± 135.7 days, respectively. Multivariable analysis showed significant risk factors of the sICH, such as warfarin over NoACs; hypertension; diabetes mellitus; brain tumors; and decreased duration of oral anticoagulation. NoACs demonstrated a lower risk of sICH than warfarin in a real-world setting using OMOP CDM confined to a single institution. Clinical studies using a CDM for the multicenter datasets may provide more reliable information about the risk of sICH.







Congratulations to the team of Young-Eun Kwon, Shin-Young Ahn, Gang-Jee Ko, Young-Joo Kwon and Ji-Eun Kim on the publication of **Impact of Uric Acid Levels on Mortality and Cardiovascular Outcomes in Relation to Kidney** Function in the Journal of Clinical Neuroscience.





Article

Impact of Uric Acid Levels on Mortality and Cardiovascular Outcomes in Relation to Kidney Function

Young-Eun Kwon 1, Shin-Young Ahn 1,2 , Gang-Jee Ko 1,2, Young-Joo Kwon 1,2 and Ji-Eun Kim 1,2,*

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Abstract: Background: Uric acid levels are linked to cardiovascular outcomes and mortality, especially in chronic kidney disease (CKD). However, their impact across varying kidney function remains unclear. Methods: We conducted a retrospective cohort study using the Observational Medical Outcomes Partnership Common Data Model (OMOP-CDM) database from a single center. Adult patients with at least one serum uric acid measurement between 2002 and 2021 were included and categorized by estimated glomerular filtration rate (eGFR): normal kidney function (≥90 mL/min/1.73 m²), mild dysfunction (60-89 mL/min/1.73 m²), moderate dysfunction (30-59 mL/min/1.73 m²), and advanced dysfunction (<30 mL/min/1.73 m²). The primary outcome was all-cause mortality with secondary outcomes being myocardial infarction (MI) and heart failure (HF). Results: A total of 242,793 participants were analyzed. Uric acid levels showed a U-shaped association with all-cause mortality in advanced kidney dysfunction, where both low (<3 mg/dL) and high (>10 mg/dL) levels increased mortality risk. In mild kidney dysfunction, lower uric acid levels were linked to better survival. HF risk increased linearly with higher uric acid, particularly in normal kidney function, while no significant association was found between uric acid and MI in any group. Conclusions: Uric acid levels are associated with mortality in a U-shaped pattern for advanced kidney dysfunction, while lower levels appear protective in mild dysfunction. These findings suggest the need for personalized uric acid management in CKD patients based on their kidney function.

Keywords: uric acid; kidney dysfunction; chronic kidney disease; mortality; cardiovascular



Academic Editor: Laetitia Dou Received: 8 October 2024 Revised: 5 December 2024









Congratulations to the team of Mitchell Conover, Patrick Ryan, Yong Chen, Marc Suchard, George Hripcsak, and Martijn Schuemie on the publication of **Objective study** validity diagnostics: a framework requiring pre-specified, empirical verification to increase trust in the reliability of real-world evidence in JAMIA.

Journal of the American Medical Informatics Association, 2025, 1–8 https://doi.org/10.1093/jamia/ocae317



Research and Applications

Objective study validity diagnostics: a framework requiring pre-specified, empirical verification to increase trust in the reliability of real-world evidence

Mitchell M. Conover , PhD^{1,2,*}, Patrick B. Ryan, PhD^{1,2,3}, Yong Chen , PhD^{1,4}, Marc A. Suchard , MD, PhD^{1,5,6}, George Hripcsak, MD^{1,3}, Martijn J. Schuemie, PhD^{1,2,5}

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*Corresponding author: Mitchell M. Conover, PhD, Observational Health Data Analytics, Johnson & Johnson, 920 US-202, Raritan, NJ 08869, United States (mconove1@its.jnj.com)

Abstract

Objective: Propose a framework to empirically evaluate and report validity of findings from observational studies using pre-specified objective diagnostics, increasing trust in real-world evidence (RWE).

Materials and Methods: The framework employs objective diagnostic measures to assess the appropriateness of study designs, analytic assumptions, and threats to validity in generating reliable evidence addressing causal questions. Diagnostic evaluations should be interpreted before the unblinding of study results or, alternatively, only unblind results from analyses that pass pre-specified thresholds. We provide a conceptual overview of objective diagnostic measures and demonstrate their impact on the validity of RWE from a large-scale comparative new-user study of various antihypertensive medications. We evaluated expected absolute systematic error (EASE) before and after applying diagnostic thresholds, using a large set of negative control outcomes.

Results: Applying objective diagnostics reduces bias and improves evidence reliability in observational studies. Among 11 716 analyses (EASE– = 0.38), 13.9% met pre-specified diagnostic thresholds which reduced EASE to zero. Objective diagnostics provide a comprehensive and empirical set of tests that increase confidence when passed and raise doubts when failed.

Discussion: The increasing use of real-world data presents a scientific opportunity; however, the complexity of the evidence generation process poses challenges for understanding study validity and trusting RWE. Deploying objective diagnostics is crucial to reducing bias and improving reliability in RWE generation. Under ideal conditions, multiple study designs pass diagnostics and generate consistent results, deepening understanding of causal relationships. Open-source, standardized programs can facilitate implementation of diagnostic analyses.

Conclusion: Objective diagnostics are a valuable addition to the RWE generation process.

Key words: observational study; research design; data interpretation, statistical; methods; causality.





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







Upcoming Workgroup Calls



Date	Time (ET)	Meeting		
Tuesday	12 pm	Generative AI and Analytics		
Wednesday	8 am	Psychiatry		
Thursday	8 am	India Community Call		
Thursday	10 am	Themis		
Thursday	11 am	Industry		
Thursday	12 pm	HADES		
Thursday	7 pm	Dentistry		
Friday	10 am	GIS-Geographic Information System		
Friday	10:30 am	Open-Source Community		
Friday	11:30 am	Steering		
Friday	11:30 am	Clinical Trials		
Monday	10 am	Getting Started (Healthcare Systems sub-workgroup)		
Monday	10 am	Africa Chapter		
Monday	11 am	Data Bricks User Group		
Monday	2 pm	Electronic Animal Health Records		



Next CBER Best Seminar: Jan. 15

Topic: Emulation of Target Trial on Vaccinations During Pregnancy

Presenter: Sonia Hernández-Díaz, MD, DrPH, Professor of Epidemiology, Harvard T.H. Chan School of Public Health

Date/Time: Jan. 15, 11 am ET



ohdsi.org/cber-best-seminar-series





Guideline-Driven Opportunities

Guideline-driven evidence generation opportunities

1d

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Chungsoo_Kim

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Treating obesity is receiving tremendous attention these days. The attached links are clinical guidelines for obesity management.

2013 AHA/ACC/TOS Guideline for the Management of Overweight and Obesity in Adults 2

Ob 202

Thamir

Thanks @Patrick_Ryan , for discussing this very interesting and bright idea. I go with what @Chungsoo_Kim mentioned especially the new standard of care of care guidelines (2025)

https://diabetesjournals.org/care/article/48/Supplement_1/S6/157564/Summary-of-RevisionsHot Standards-of-Care-in-Diabetes . In these guidelines, there were some updated recommendations for the use of new dual GIP and GLP medications. Also, the role of SGLT2 inhibitors in MACE and HF. These might be discussed in the LEGEND studies but also can be studied further

especially comparing combinations with mono therapy, or event safety profile.



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Another area I guess it is not touched much is the depression treatments, a recent guideline Canadian Network for Mood and Anxiety Treatments (CANMAT) 2023 Update on Clinical Guidelines for Management of Major Depressive Disorder in Adults discussed the different medications for depression. Treatment Guidelines: Depression | aapp.org

As you know, these medications have benefits but not sure if there were any comparison between different groups like SSRI. TCAs. or SNRI or others. Also, these medications have

between different groups like **SSRI**, **TCAs**, or **SNRI** or others. Also, these medications have big concerns about their safety that also can be discussed. I am not really expert in these areas and I am sure there are who know better than me, but I thought I'd open this up for future discussion.

We're currently in the process of developing analysis tools that address these gaps in specific disease areas and treatment stages.

We'd love to hear your thoughts!

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further complicating the decision-making process.

Cancer guidelines are often large, intricate, and subject to frequent updates due to multifaceted nature of the disease. Cancer is not a single disease but a collection of hundreds of different conditions, each with its own biology, behavior, and response to treatment. This requires guidelines to account for a wide range of variables such as tumor type, stage, genetic mutations, and patient characteristics. Additionally, cancer treatment often involves multidisciplinary approaches, including surgery, radiation, chemotherapy, immunotherapy, and targeted therapies,

Several organizations publish guidelines but let us focus on NCCN guidelines, these are step-bystep decision trees for each cancer type all the way down the possible decision points. Unfortunately, the guidelines are much less specific than one would expect. It's full of sentences like:

- "With the recent changes to first-line treatment options for metastatic disease, many
 providers are moving towards immune checkpoint inhibitor combinations, such as
 enfortumab vedotin plus pembrolizumab, as a first-line treatment option. In this evolving
 paradigm, there is limited evidence to guide the optimal selection of second- and
 subsequent-line therapies following these new first-line regimens." NCCN Bladder
 Cancer
- "Clinical trial enrollment is recommended by the NCCN Panel for all patients when appropriate, but is strongly recommended for second-line and subsequent therapies, as data for locally advanced or metastatic disease treated with subsequent-line therapy are highly variable." — NCCN Bladder Cancer
- "The optimal sequencing of systemic therapy and resection remains unclear." NCCN Colon Cancer

Bottom line: We have large and complicated decision trees, somewhat fuzzy, with each node its own constellation. Most of them are based on clinical trials. With respect to RWE, we could create evidence for each node about:

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Workgroup OKRs

Each year, workgroup representatives join a February community call to present the mission, objectives and key results for their respective groups. These 2-4 minute presentations are recorded and posted on the Workgroups homepage on OHDSI.org.

Please choose a date to sign up for a February date; once a date has at least 10 workgroups, it will be closed.





The Center for Advanced Healthcare Research Informatics (CAHRI) at Tufts Medicine welcomes:



Vipina Keloth, PhD

Associate Research Scientist in Biomedical Informatics and Data Science at Yale University School of Medicine

'Exploring the realm of large language models for information extraction in the biomedical domain'

January 23, 2025, 11am-12pm EST Virtually via **Zoom**





CDM Survey Subgroup Landscape Assessment

The CDM Survey Subgroup invites colleagues who have or are going to design, develop, and/or implement research surveys and use them with the OMOP CDM to share information about those efforts by completing this survey. Your completion of this 10-15 minute survey will provide information to the CDM workgroup about OMOP utilization among survey research teams. The CDM Survey subgroup is a collaborative effort, led by a team at the National Cancer Institute, to develop standardized approaches and best practices for helping research teams better integrate survey data elements into the OMOP common data model.

The deadline has been extended to mid-January.

LANDSCAPE ASSESSMENT

Activities

- Invite representatives from cohorts with experience using the CDM for survey data to share their knowledge and challenges.
- Conduct a community survey to gather information on experiences and needs related to survey data in the CDM.
- Review the most used Common Data Elements (CDMs) as a foundation for developing standards, tools, and best practices.

Key Result

 A comprehensive report summarizing survey CDM mapping resources, challenges, and identified development priorities (vocabulary, standards, tools, best practices) to be shared with the OHDSI community.

WHO SHOULD PARTICIPATE

- You have survey data and you've mapped it to the OMOP CDM
- You have survey data and you would like to map it to the OMOP CDM
- You are in the process of developing a survey(s) and plan to map to the OMOP CDM
- Multiple perspectives from the same team
- Multiple surveys from the same person





Monday

Lessons from mapping cancer information from European hospitals to ICD-O-3 conditions in OMOP

(Lars Halvorsen, Olivier Bouissou, Elisabeth Ross, Stelios Theophanous, Joëlle Thonnard, Piers Mahon)

Lessons from mapping cancer information from European hospitals to ICD-O-3 conditions in OMOP

♣ PRESENTER: Lars Halvorsen

- · European hospitals in the DigiONE network¹ face a recurring conversion challenge in mapping cancer conditions.
- The hospitals mapped their cancer diagnoses to ICD-O-3 or SNOMED using topography and morphology information from pathology reports or electronic healthcare records.
- Differences in terminologies used at source presented different challenge for different hospitals.
- A sizeable subset of ICD-O-3 diagnosis codes were found to not vet be defined in the OHDSI vocabularies2

Different paths for mappings to ICD-O-3 condition concepts:

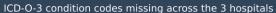
· ICD-O-3 condition codes derived from ICD-O-3 morphology and topography codes.

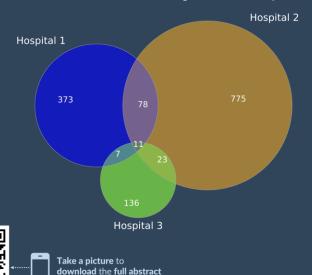
- · Pathology results coded in terminology based on old SNOMED
- · Most T codes correspond to current SNOMED concept, while most M codes had equivalent ICD-O-3 morphology code
- · These T and M codes were combined to ICD-O-3 condition concepts, with some exceptions needing special mapping

Hospital 3:

- · ICD-O-3 topography and ICD-O-3 morphology codes available in the source system
- · Direct construction of ICD-O-3 condition codes.

Including additional ICD-O-3 condition concepts in the OHDSI vocabularies will facilitate cancer research across hospitals

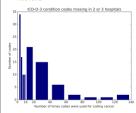




· All 3 hospitals had ICD-O-3 condition codes not yet available in the OHDSI vocabularies. 119 such missing codes appeared in at least two of the three hospitals (see abstract for full list).

Solid and harmatological concers OHDSI vocabulary version		Hospital 1 v5.0 29-FEB-24	Wospital 2 v5.0 23-JAN-23	Hospital 3 v5.0 29 FEB-24
	# unique source codes	3588	2979	1823
Mapping Result	# standard ICD-O-3 Condition concept ID	2413	1524	1306
	# non-standard ICD-O-3 Condition concept IDs mapped to standard (\$NOMED) concept IDs	706	568	338
	# ICD-O-3 Condition codes with no concept ID in vocabulary	469	887	177

- Of these missing codes, 53 have already been identified for inclusion in a future OHDSI vocabulary release, while 66 of these codes had not yet been included.
- The DigiONE hospitals will further evaluate these findings and work with the Oncology WG - Vocabulary subgroup to include an additional set of ICD-O-3 condition codes considered as



The majority of codes missing in 2 or 3 hospitals were rarely used for coding cancer, with 71 codes used less than 10 times. Only a few codes were used more frequently, with 4 codes used between 80 and 140

References

- Authors: Lars Halvorsen, Olivier Bouissou, Elisabeth Ross, Stelios Theophanous, Joëlle Thonnard, Piers Mahon













Tuesday

Jackalope Plus Performance: Benchmarking and Competitors

(Denys Kaduk, Bohdan Khilchevskyi, Maksym Trofymenko, Tetiana Nesmiian, Polina Talapova, Max Ved, Inna Ageeva)

Jackalope Plus Performance: Benchmarking and Competitors



PRESENTER:
Polina Talapova

polina.talapova @sciforce.tech

INTPO

In the healthcare domain, where data plays an increasingly crucial role, accurate and efficient mapping of medical terminologies is fundamental. This mapping ensures seamless data exchange (interoperability) between different healthcare information systems, facilitates clinical research efforts, and ultimately improves patient care. The advent of the Observational Medical Outcomes Partnership (OMOP) Common Data Model (CDM) has further emphasized the significance of terminology mapping.

While manual mapping remains an option, it is a laborious and time-consuming process, hindering large-scale data standardization initiatives. To address this challenge, several automated mapping tools have emerged, aiming to streamline the process and enhance efficiency. This study benchmarks Jackalope Plus, a prominent mapping tool, against manual mapping and Usagi, another automated option, to assess their relative strengths and weaknesses.

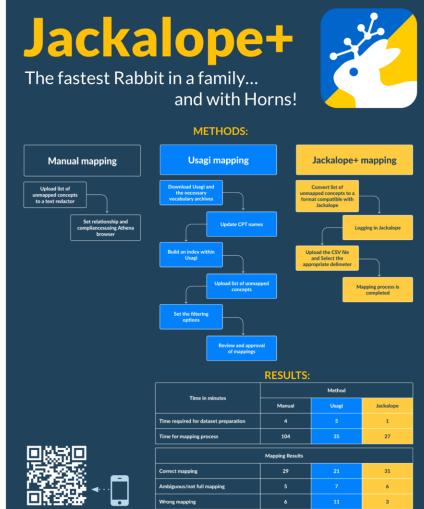
METHOD

Three domain experts (testers) familiar with all mapping methods and assigned tasks based on their preferences. The benchmark utilized a newly created dataset comprising unmapped concepts from MeSH and CIEL, as well as some concepts created by our team expert. The test set was provided to the testers prior to the mapping process, formatted as a CSV file with a semicolon (;) delimiter containing 40 unmapped concepts.

Three methods were used:

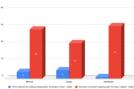
- 1. Manual mapping
- 2. Usagi mapping
- 3. Jackalope Plus mapping

Each mapping method followed specific steps, described at the center of the poster..





RESULTS



Results are presented in the left lower corner and briefly on a diagram above. The time for preparation is not included in the final results due to the variability of the source data.

The benchmarking process revealed significant insights into the efficiency and accuracy of the mapping tools compared. Both Jackalope Plus and Usagi demonstrated the ability to significantly reduce the time required for the mapping process compared to manual methods. This time efficiency is critical in large-scale data standardization efforts, where speed and accuracy are paramounts.

However, the quality of the results varied between the tools. Jackalope Plus consistently produced more accurate mappings than Usagi. This superior accuracy can be attributed to Jackalope Plus's advanced algorithms and streamlined workflow, which minimize errors and enhance precision. The domain experts reported that Jackalope Plus not only reduced the time spent on mapping but also delivered higher-quality results, making it a more reliable tool for integrating medical data into standardized frameworks.

THETEAN

Denys Kaduk, Bohdan Khilchevskyi, Maksym Trofymenko, Tetiana Nesmiian, Polina Talapova, Max Ved, Inna Ageeva









Wednesday

Trade-offs in the design of explainable prediction models for health care

(Aniek F. Markus, Jan A. Kors, Katia M.C. Verhamme, Peter R. Rijnbeek) "The human body is a black box", do we need Explainable AI (XAI)?

Trade-offs in the design of explainable prediction models for health care

Background: Artificial intelligence (AI) has the potential to improve patient care, but implementation of prediction models in clinical practice is still limited. Lack of transparency is - at least in the current state of AI maturity often seen as one of the main problems. This work explores different types of explanations to overcome the transparency problem of AI in health care.





Empirical research using OMOP data

Using the PLP framework, we applied different types of XAI techniques to real-world data, which led to the following findings: - Interpretable models with a limited number of covariates and good predictive performance can be developed for various prediction tasks (e.g. using clinical expertise, feature selection or rule-based

- Model are unstable both in terms of the variables included in the model and in the sign of their coefficients. Similarly, different feature importance methods result in different generated explanations. - There is some trade-off between model performance and

interpretability, but it varies across prediction tasks and seems to be stronger for high levels of model complexity

Identified risks of XAI

- Often multiple explanations possible (e.g. model instability, feature importance disagreement).
- 2 Explanations can be overinterpreted (e.g. as causal elation, to identify risk factors).
- Requiring (certain types of) explanations might come at cost of predictive performance.
- Explanations can have unintended (adverse) effects (e.g. decreasing human-machine task performance)

Conclusion: Although explanations can be useful to assist implementation in practice by allowing for a human in the loop to detect and correct problems (e.g. existing biases), explanations are never sufficient by itself and not the ultimate goal. It is important to link the need an explanation strives to fulfil with the design choice.







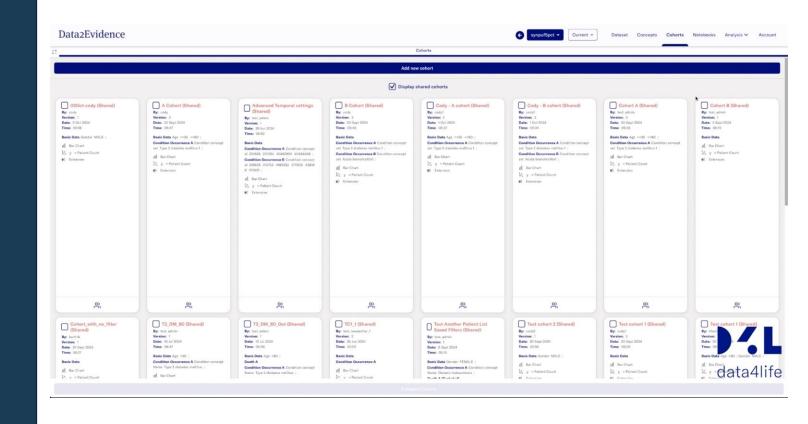
in ohdsi



Thursday

An interactive approach for data exploration and phenotyping in the Data2Evidence platform

(Satish Anbazhagan, Peter Hoffmann)

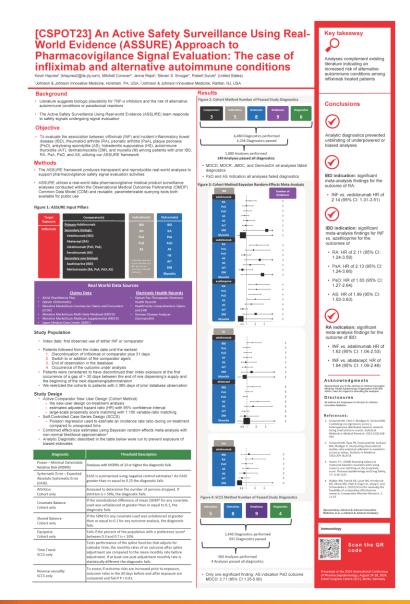




Friday

An Active Safety Surveillance
Using Real-World Evidence
(ASSURE) Approach to
Pharmacovigilance Signal
Evaluation: The case of
infliximab and alternative
autoimmune conditions

(Kevin Haynes, Mitchell M. Conover, Jenna Reps, Steven S. Smugar, Robert Suruki)





Where Are We Going?

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



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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



