

Establishing scientific best practices for real-world analysis

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How to achieve reliable evidence?

• **Transparency**: others should be able to reproduce your study in every detail using the information you provide.

• **Prespecify** what you're going to estimate and how, to avoid hidden multiple testing (fishing expeditions, p-value hacking). Run your analysis only once.

• Validation of your analysis: you should have evidence that your analysis does what you say it does, and that what it does is the right way to answer the question.



How to achieve reliable evidence?

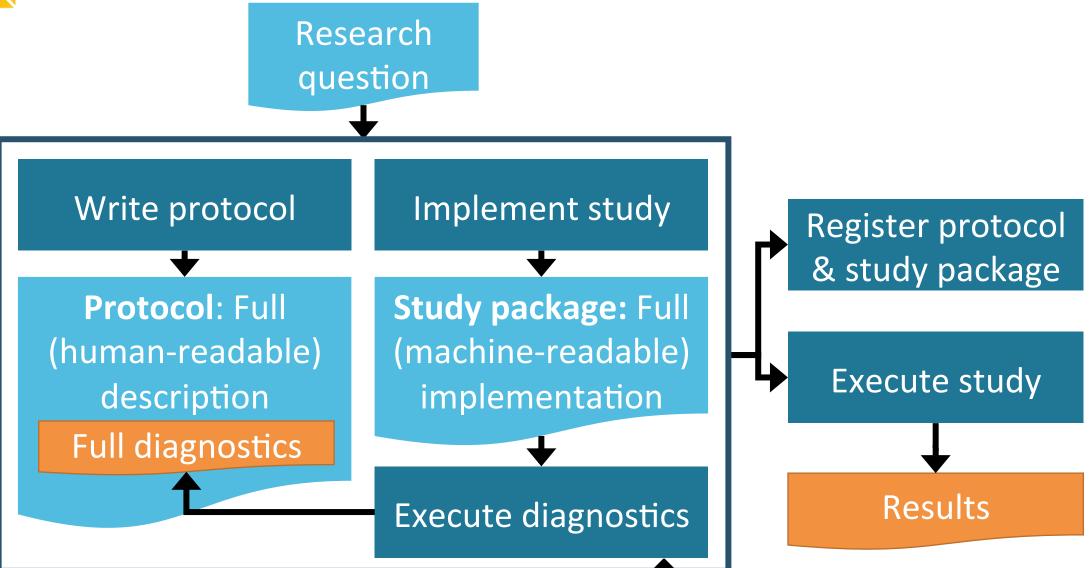
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Transparency & Prespecification



Without looking at the outcome of interest!



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Study validity

Evidence Quality



Data Quality: Are the data of sufficient quality for our research purposes?



Clinical Validity: To what extent does the analysis conducted match the clinical intention?



Software Validity: Does the software do what it is expected to do?



Method Validity: Is this method valid for answering this question?





Are the data of sufficient quality for our research purposes?

Just discussed by Clair and Andrew:



DATA QUALITY ASSESSMENT

SYNTHEA SYNTHETIC HEALTH DATABASE

Results generated at 2019-08-21 18:08:28 in 13 mins

	Verification				Validation				Total			
	Pass	Fail	Total	% Pass	Pass	Fail	Total	% Pass	Pass	Fail	Total	% Pass
Plausibility	159	21	180	88%	283	0	283	100%	442	21	463	95%
Conformance	637	34	671	95%	104	0	104	100%	741	34	775	96%
Completeness	369	17	386	96%	5	10	15	33%	374	27	401	93%
Total	1165	72	1237	94%	392	10	402	98%	1557	82	1639	95%





Are the data of sufficient quality for our research purposes?

Just discussed by Clair and Andrew:



DATA QUALITY ASSESSMENT

EDM Forum Community

eGEMs (Generating Evidence & Methods to improve patient outcomes)

Publish

9-11-2016

A Harmonized Data Quality Assessment Terminology and Framework for the Secondary Use of Electronic Health Record Data

Michael G. Kahn
University of Colorado Anschutz Medical Campus, michael.kahn@ucdenver.edu

PATABASE

3:28 in 13 mins

⁄er	ification			Va	lidation		Total				
il	Total	% Pass	Pass	Fail	Total	% Pass	Pass	Fail	Total	% Pass	
1	180	88%	283	0	283	100%	442	21	463	95%	
4	671	95%	104	0	104	100%	741	34	775	96%	
7	386	96%	5	10	15	33%	374	27	401	93%	
2	1237	94%	392	10	402	98%	1557	82	1639	95%	





To what extent does the analysis conducted match the clinical intention?

For example, will we answer

"Do ACE inhibitors cause angioedema"

or

"Do ACE inhibitors cause suspicion of angioedema"



Clinical Validity

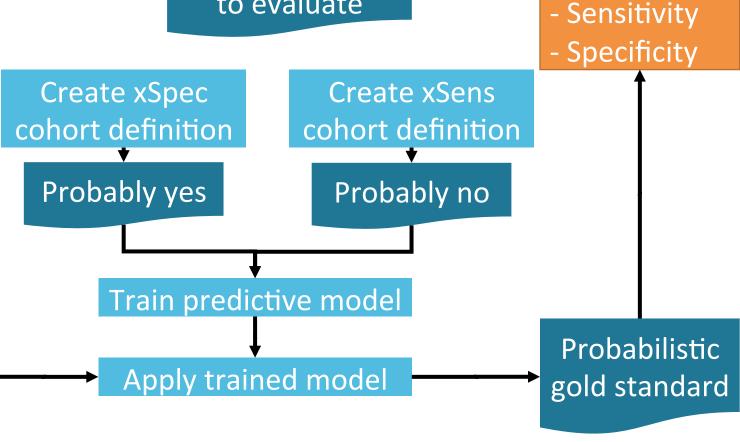
To what extent does the analysis conducted match the clinical intention?

Large

random

sample

- Chart review
- PheValuator



Cohort definition

to evaluate

Evaluate

- PPV





To what extent does the analysis conducted match the clinical intention?

Cohort definition to evaluate

Evaluate

- PPV
- Sensitivity
- Specificity



Contents lists available at ScienceDirect

Journal of Biomedical Informatics

journal homepage: www.elsevier.com/locate/yjbin



Create xSens hort definition

Probably no

model

odel



PheValuator: Development and evaluation of a phenotype algorithm evaluator

evaluator

Joel N. Swerdel^{a,b,*}, George Hripcsak^{b,c}, Patrick B. Ryan^{a,b,c}

- ^a Janssen Research & Development, 920 Route 202, Raritan, NJ 08869, USA
- b OHDSI Collaborators, Observational Health Data Sciences and Informatics (OHDSI), 622 West 168th Street, PH-20, New York, NY 10032, USA
- ^c Columbia University, 622 West 168th Street, PH20, New York, NY 10032, USA

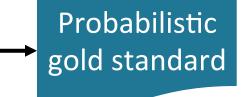


Keywords:
Phenotype algorithms
Validation
Diagnostic predictive modeling

ABSTRACT

Background: The primary approach for defining disease in observational healthcare databases is to construct phenotype algorithms (PAs), rule-based heuristics predicated on the presence, absence, and temporal logic of clinical observations. However, a complete evaluation of PAs, i.e., determining sensitivity, specificity, and positive predictive value (PPV), is rarely performed. In this study, we propose a tool (PheValuator) to efficiently estimate a complete PA evaluation.

Methods: We used 4 administrative claims datasets: OptumInsight's de-identified Clinformatics™ Datamart (Eden



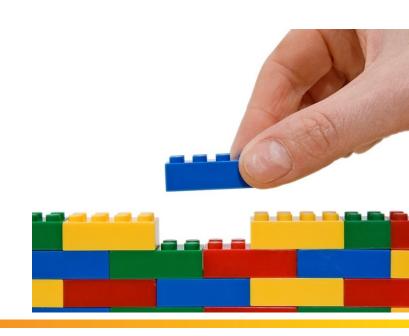




Does the software do what it is expected to do?

- Building blocks: OHDSI Standardized Tools
 - Unit tests
 - Simulations

- Configuration of blocks: Study package
 - Code review
 - Double coding









Method Validity

Is this method valid for answering this question?

This requires a method that can be validated!

A **systematic** approach, for example our large-scale propensity models:

- Create a large set of covariates (10,000 < n 100,000)
- Use LASSO to fit propensity model
- Match / stratify on propensity score
- Check that covariate balance is achieve on all observed variables





This requires a method that can be validated!

Drug Saf DOI 10.1007/s40264-017-0581-7 ORIGINAL RESEARCH ARTICLE

Channeling in the Use of Nonprescription Paracetamol and Ibuprofen in an Electronic Medical Records Database: Evidence and Implications

Rachel B. Weinstein¹ • Patrick Ryan¹ • Jesse A. Berlin² • Amy Matcho³ • Martijn Schuemie¹ • Joel Swerdel¹ • Kayur Patel⁴ • Daniel Fife¹

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Abstract

Introduction Over-the-counter analgesics such as paracetamol and ibuprofen are among the most widely used, and having a good understanding of their safety profile is

distributions between drugs, and examined the degree to which channeling bias could be controlled using a combination of negative control disease outcome models and large-scale propensity score matching. Analyses were



International Journal of Epidemiology, 2018, 1–10 doi: 10.1093/ije/dyy120 Original article



Original article

Evaluating large-scale propensity score performance through real-world and synthetic data experiments

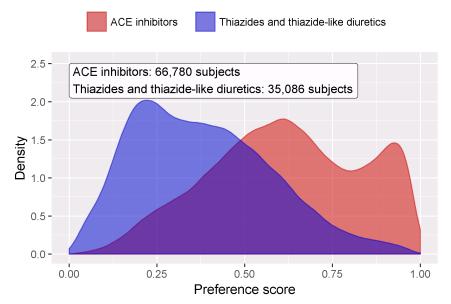
Yuxi Tian, 1* Martijn J Schuemie 2 and Marc A Suchard 1,3,4

¹Department of Biomathematics, David Geffen School of Medicine at UCLA, University of California, Los Angeles, CA, USA, ²Epidemiology Department, Janssen Research and Development LLC, Titusville, NJ, USA, ³Department of Biostatistics, UCLA Fielding School of Public Health, University of California, Los Angeles, CA, USA and ⁴Department of Human Genetics, David Geffen School of Medicine at UCLA, University of California, Los Angeles, CA, USA

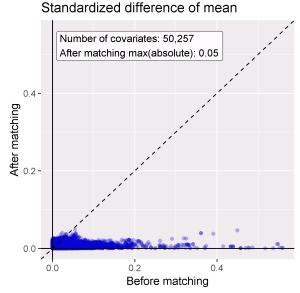




• Do we pass our (design-specific) study diagnostics? E.g.



Are the exposure groups sufficiently comparable?

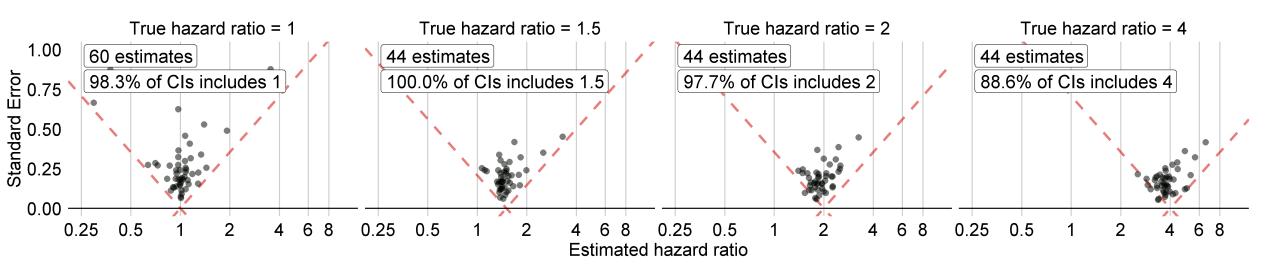


Do we balance on all observed variables?





- Do we pass our (design-specific) study diagnostics?
- Do negative and positive controls show nominal operating characteristics? (after empirical calibration)







Do we pass our (design-specific) study diagnostics?



hal operating

Empirical confidence interval calibration for population-level effect estimation studies in observational healthcare data

Martijn J. Schuemie^{a,b,1}, George Hripcsak^{a,c,d}, Patrick B. Ryan^{a,b,c}, David Madigan^{a,e}, and Marc A. Suchard^{a,f,g,h}

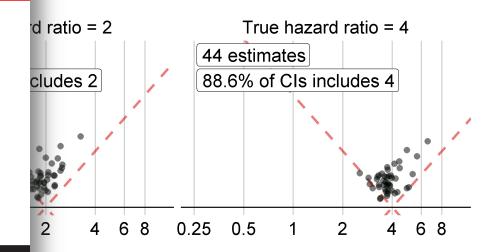
Observational Health Data Sciences and Informatics, New York, NY 10032; Epidemiology Analytics, Janssen Research & Development, Titusville, NJ 08560; Department of Biomedical Informatics, Columbia University, New York, NY 10032; Medical Informatics Services, New York-Presbyterian Hospital, New York, NY 10032; Department of Statistics, Columbia University, New York, NY 10027; Department of Biomathematics, University of California, Los Angeles, CA 90095; ⁹Department of Biostatistics, University of California, Los Angeles, CA 90095; and ^hDepartment of Human Genetics, University of California,

Edited by Victoria Stodden, University of Illinois at Urbana-Champaign, Champaign, IL, and accepted by Editorial Board Member Susan T. Fiske October 26, 2017 (received for review June 15, 2017)

Observational healthcare data, such as electronic health records and administrative claims, offer potential to estimate effects of medical products at scale. Observational studies have often been found to be nonreproducible, however, generating conflicting results even when using the same database to answer the same question. One source of discrepancies is error, both random caused by sampling variability and systematic (for example, because of confounding, selection bias, and measurement error). Only random error is typically quantified but converges to zero as databases become larger, whereas systematic error persists independent from sample size and therefore, increases in relative importance. Negative controls are exposure-outcome pairs,

age treatment effect. Systematic error can manifest from multiple sources, including confounding, selection bias, and measurement error. While there is widespread awareness of the potential for systematic error in observational studies and a large body of research that examines how to diagnose and statistically adjust for specific sources of bias, there has been comparatively little work in devising approaches to empirically estimate the magnitude of systematic error or clinical applications that show how to integrate this error into effect estimation methods.

The acuity of this problem is only exacerbated as the size of observational databases grow: random error (the only component that is typically quantified) converges to zero as sample



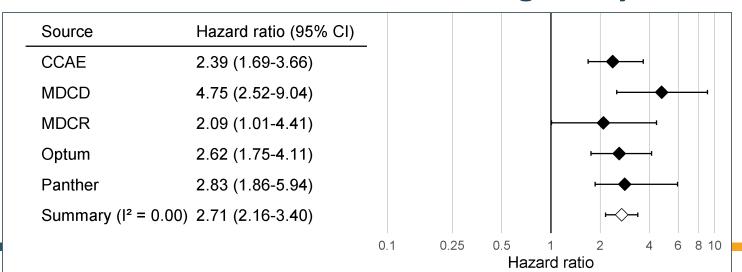
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- Do we pass our (design-specific) study diagnostics?
- Do negative and positive controls show nominal operating characteristics?
- Do we observe between-database heterogeneity?







Do we pass our (design-specific) study diagnostics?



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DOI: 10.1093/aje/kwt010
Advance Access publication:

minal operating

Practice of Epidemiology

Evaluating the Impact of Database Heterogeneity on Observational Study Results

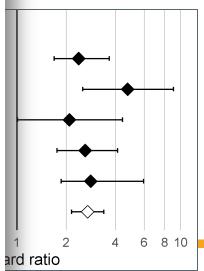
David Madigan*, Patrick B. Ryan, Martijn Schuemie, Paul E. Stang, J. Marc Overhage, Abraham G. Hartzema, Marc A. Suchard, William DuMouchel, and Jesse A. Berlin

 * Correspondence to Dr. David Madigan, Department of Statistics, Columbia University, 1255 Amsterdam Avenue, New York, NY 10027 (e-mail: david.madigan@columbia.edu).

Initially submitted November 11, 2012; accepted for publication January 17, 2013.

Clinical studies that use observational databases to evaluate the effects of medical products have become commonplace. Such studies begin by selecting a particular database, a decision that published papers invariably report but do not discuss. Studies of the same issue in different databases, however, can and do generate different results, sometimes with strikingly different clinical implications. In this paper, we systematically study heterogene-

eneity?





Best practices checklist

- Fully prespecify your study
 - Protocol
 - Study package

Finalized when you pass all study diagnostics

Must be **posted publicly**

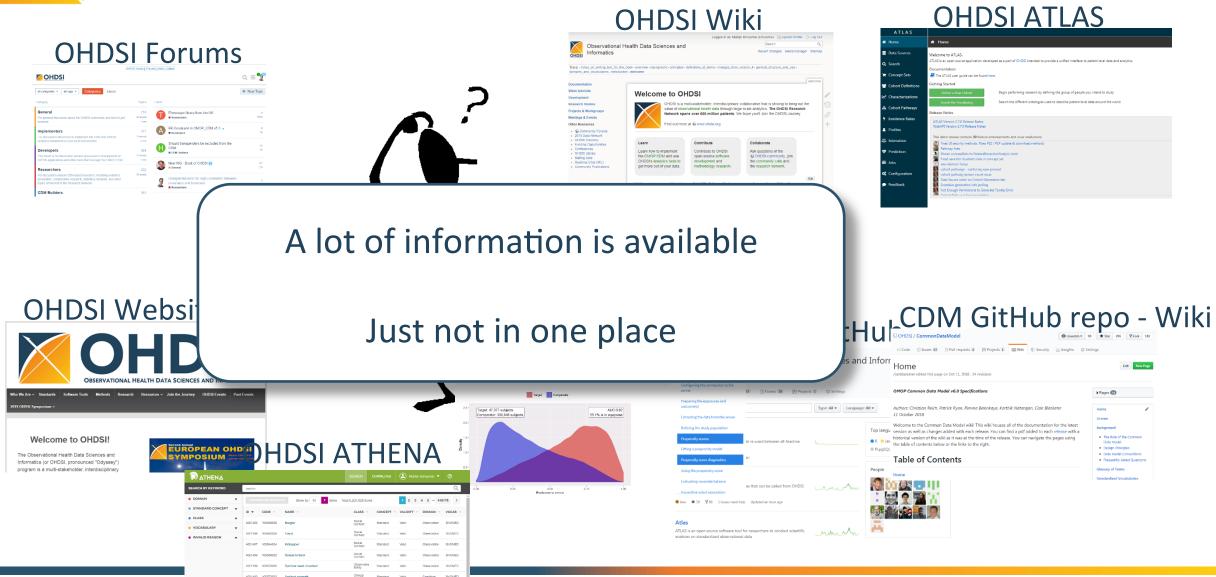
- Establish data quality using the Data Quality Dashboard and other checks
- Establish clinical validity, e.g. using PheValuator
- Establish software validity by
 - Using the OHDSI's standardized analytics tools for the standardized parts, and
 - Using code review or double coding for study-specific parts
- Establish method validity by
 - Study diagnostics
 - Using negative and positive controls
 - Executing on multiple databases



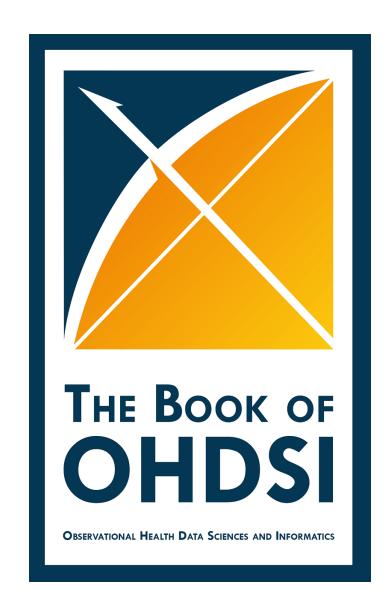
How can you follow these best practices yourself?



Where can I learn how to implement best practices?



A central knowledge repository...





- I. The OHDSI Community
- II. Uniform DataRepresentation
- III. Data Analytics
- IV. Evidence Quality
- V. OHDSI Studies



- I. The OHDSICommunity
- II. Uniform Data
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- 1. The OHDSI Community
- 2. Where to Begin
- 3. Open Science



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- 6. Extract Transform Load



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- 8. OHDSI Analytics Tools
- 9. SQL and R
- **10.Defining Cohorts**
- 11. Characterization
- 12. Population-Level Estimation
- 13. Patient-Level Prediction



Parts:

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14. Evidence Quality

15. Data Quality

16.Clinical Validity

17. Software Validity

18. Method Validity



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19.Study steps

20.OHDSI Network Research



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- Fully prespecify your study
- Establish data quality
- Establish clinical validity
- Establish software validity
- Establish method validity



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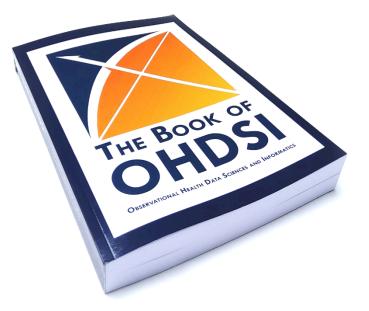
18. Method Validity

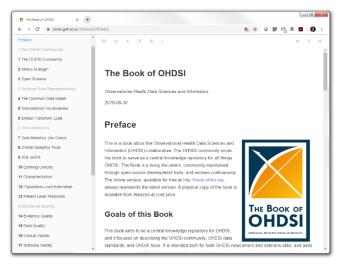
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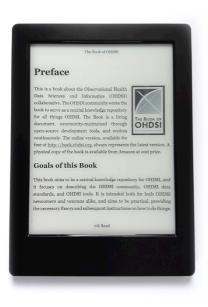
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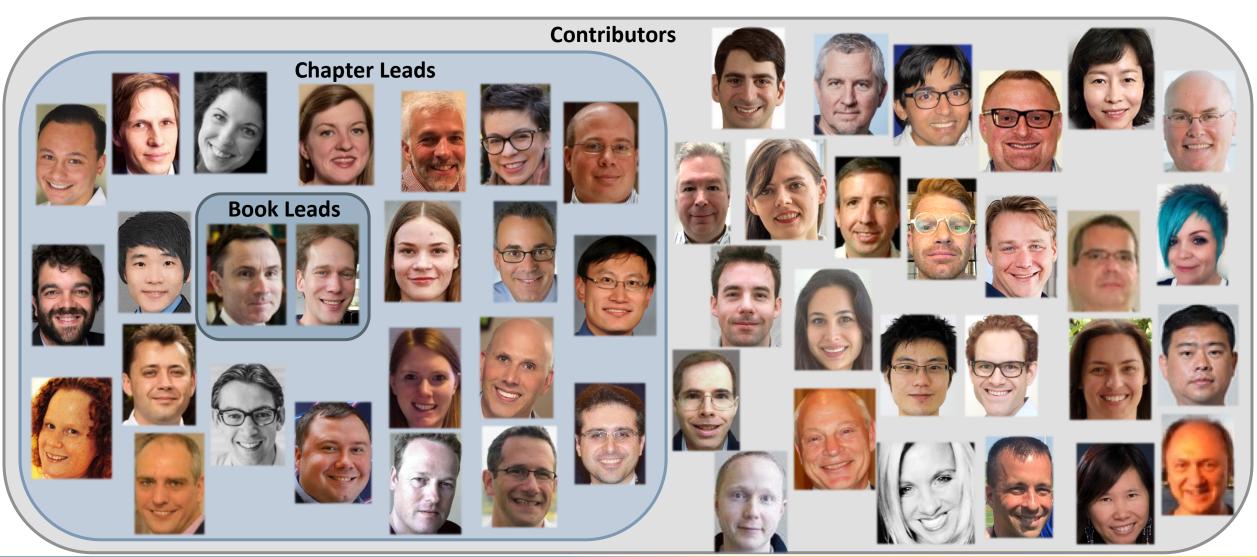
Writing a Book OHDSI-Style!

Requirements:

- Community effort
- Completely open and transparent



It takes a village...





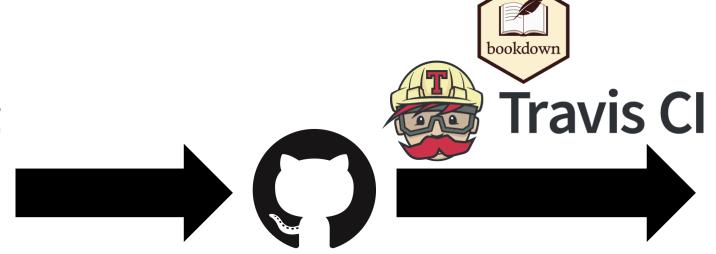
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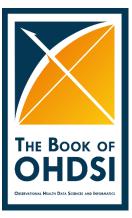




Open and Transparent Process







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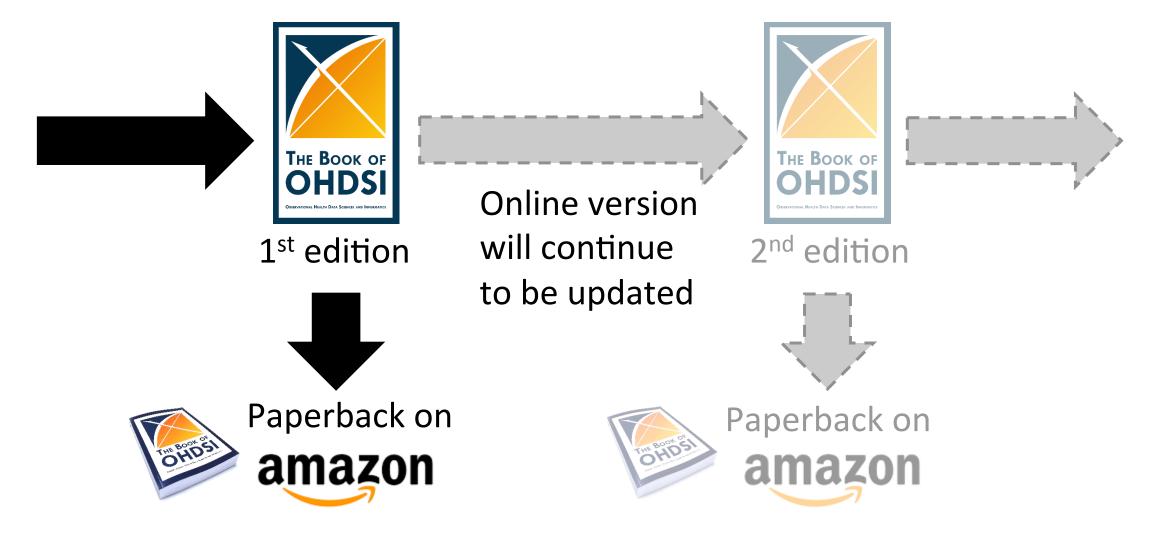
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Living Document





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