

Wrap up

David Madigan¹

Patrick Ryan²

Martijn J. Schuemie²

Marc A. Suchard³

- 1. Department of Statistics, Columbia University
- 2. Janssen Research & Development, LLC
- 3. Departments of Biomathematics and Human Genetics, David Geffen School of Medicine, and Department of Biostatistics, Fielding School of Public Health, University of California, Los Angeles



Lessons learned

 You are not getting off that easy! I am not soup-feeding a summary.

What did you learn?



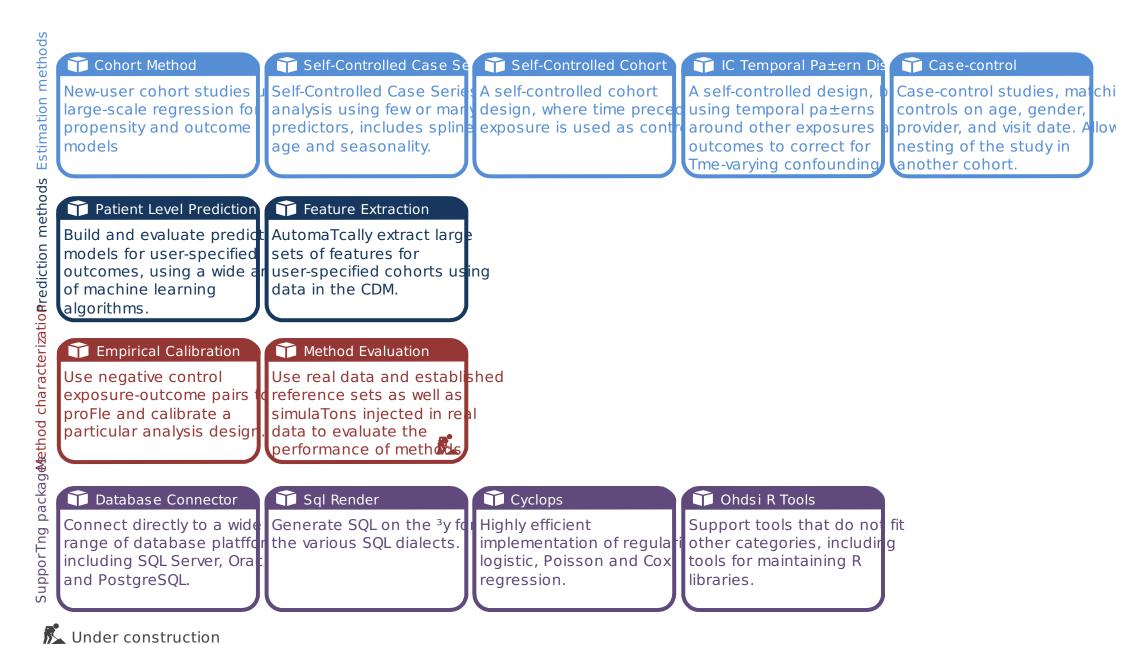
Remaining tasks

Task designers and programmers with documenting results

How to present?



OHDSI Methods library



Note other designs at scale



Infrastructure investment

- DatabaseConnector flexible interface
- SqlRender cross-SQL-platform translation and parameterization
- Cyclops fits large-scale regressions very efficiently

To support this work:

- 1. Cite!
- 2. Co-author with package developers

To develop in this sphere:

1. Contribute to the OHDSI Methods Library



Citations

To cite Cyclops in publications use:

Suchard MA, Simpson SE, Zorych I, Ryan P and Madigan D (2013). "Massive parallelization of serial inference algorithms for complex generalized linear models." ACM Transactions on Modeling and Computer Simulation, 23, pp. 10. link

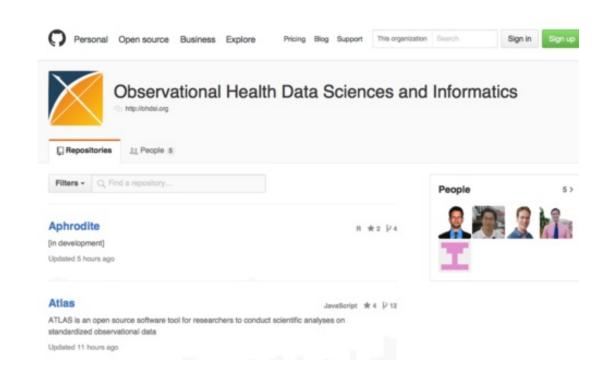
To cite CohortMethod in publications use:

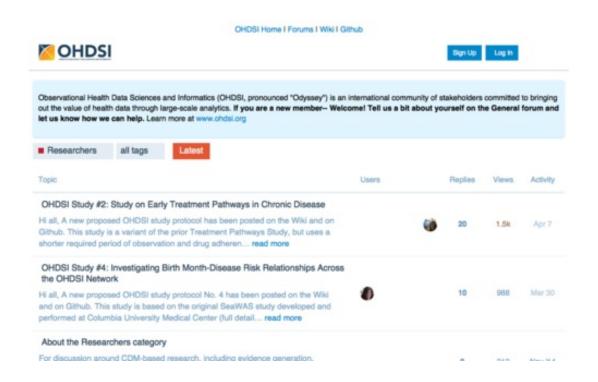
Schuemie MJ, Suchard MA and Ryan P (2015). "CohortMethod: New-user cohort method with large scale propensity and outcome models. R package version 2.0.3.



Building a community

- Community forums: Researchers tag
- Population-level estimation workgroup: github.com/OHDSI







Tutorial improvement

This course is for you:

- What went well?
- What did not?
- Welcoming feedback Survey Monkey
- Priorities for tutorials on other aspects of the R Methods Library? Other designs?



Next tutorial



Traveling short-course:

- March 2017
- Ajou Medical School, South Korea

See you there!