

# Advancing Interpretable Regression Analysis for Binary Data: A Novel Distributed Algorithm Approach

Jiayi (Jessie) Tong, Assistant Professor Department of Biostatistics, Johns Hopkins Bloomberg School of Public Health December 3, 2024

**OHDSI Community Call** 

### Background

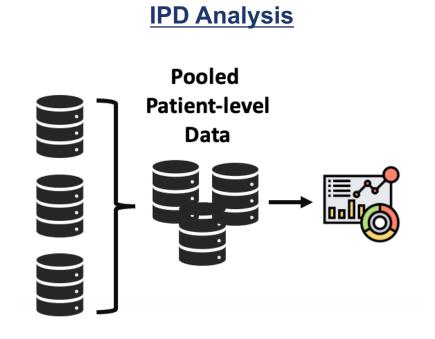
- Rare diseases -> rare binary outcome, the sparse data problem is a significant challenge.
- The lack of sufficient cases (e.g., patients with disease) in the data leads to biased estimates of the effect of a treatment or a medication in observational studies.



### Background

• Multi-site individual patient-level data (IPD) analysis increases the number of cases

 However, it is not feasible when the individual-level data from different studies cannot be shared.

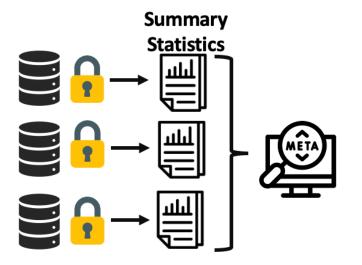




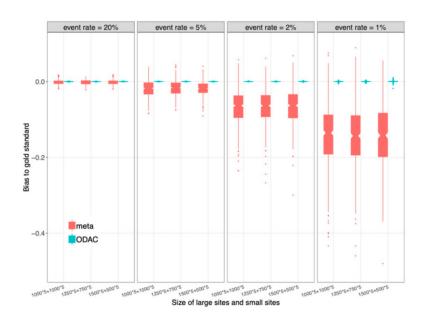
# Meta-analysis Method



# Meta-analysis (Divide and conquer)



# Meta-analysis has low accuracy for rare data



#### Federated Learning Algorithms

#### **Privacy-Preserving Federated Learning Algorithms**

- Enables fitting statistical models in a federated manner
- Requires summary statistics, instead of IPD
- Ensures data privacy and security

**Federated Learning Algorithms for Rare Binary Data** 

### Existing Federated Learning Algorithms for Binary Data

# **Logistic Regression Model -> Odds Ratio (OR)**

**Aggregated** 

**Aggregated** 

The choice between **Relative Risk (RR)** and OR has been debated in the literature, with RR being preferred in most prospective studies due to **its collapsibility** and **better interpretation**.

Poisson Regression Model -> Relative Risk (RR)

Shared Models Without Sharing data

Yuan Wu, Xiaoqian Jiang, Jihoon Kim, Lucila Ohno-Machado

Research and Applications

distribute Risk (RR)

distributed algorithm

Rui Duan (6), 1 Mary Regina Boland (6), 1 Zixuan Liu², Yue Liu, 3 Howard H Chang, 4 Hua Xu, 5 Haitao Chu, 6 Christopher H Schmid, 7 Christopher B Forrest, 8 John H Holmes, 1 Martijn J Schuemie (6), 9 Jesse A Berlin, 9 Jason H Moore, 1 and Yong Chen (6) 1



American Journal of Epidemiology Copyright © 2004 by the Johns Hopkins Bloomberg School of Public Health All rights reserved Vol. 159, No. 7 *Printed in U.S.A.* DOI: 10.1093/aje/kwh090

#### A Modified Poisson Regression Approach to Prospective Studies with Binary Data

#### **Guangyong Zou<sup>1,2</sup>**

- <sup>1</sup> Robarts Clinical Trials, Robarts Research Institute, London, Ontario, Canada.
- <sup>2</sup> Department of Epidemiology and Biostatistics, University of Western Ontario, London, Ontario, Canada.

### **Proposed Method**

#### ODAP-B: One-shot Distributed Algorithm of Modified Poisson Regression for Binary Data

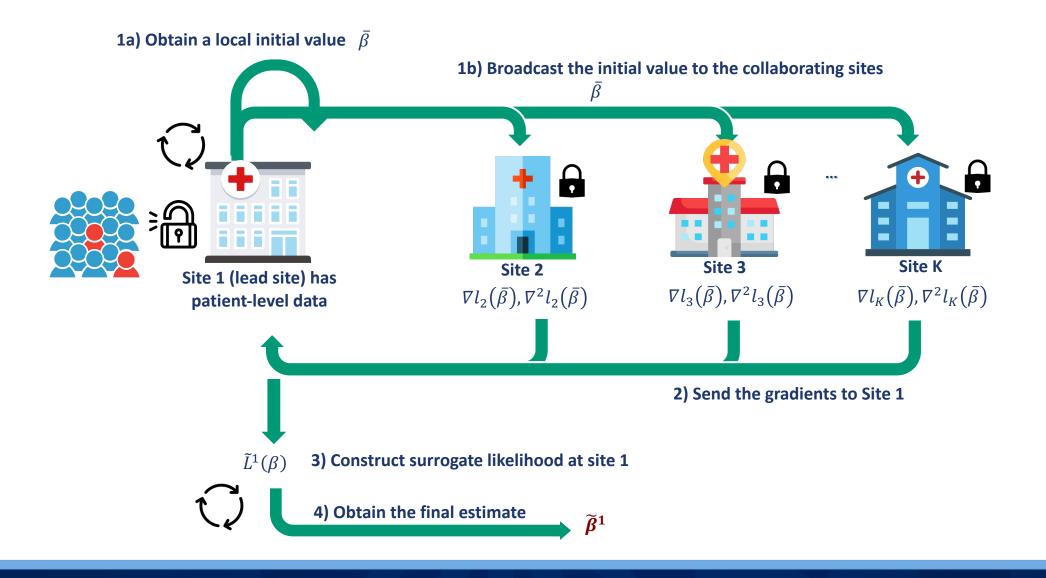
- The surrogate likelihood (SL) approach (Jordan et al 2018 JASA)
  - For some given initial value  $\bar{\beta}$ , consider Taylor expansion for the multi-site likelihood

$$L(\beta) = L(\bar{\beta}) + \nabla L(\bar{\beta})^{T} (\beta - \bar{\beta}) + \sum_{t=2}^{\infty} \frac{1}{t!} \nabla^{t} L(\bar{\beta}) (\beta - \bar{\beta})^{\otimes t}$$

For some given initial value  $\bar{\beta}$ , consider Taylor expansion for the local likelihood

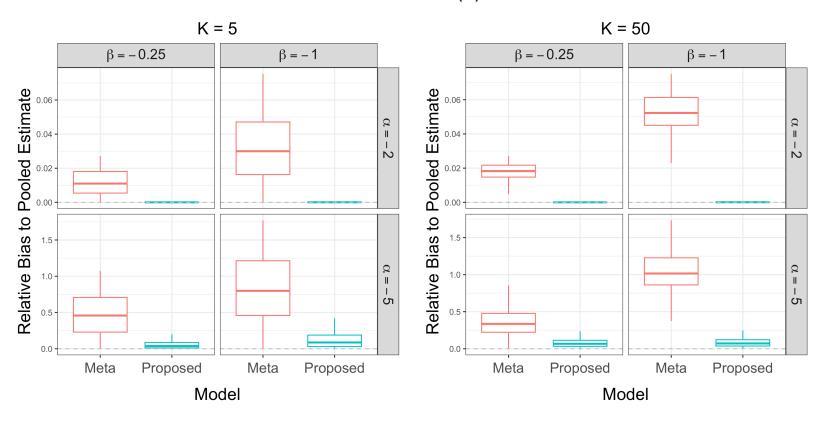
$$L_1(\beta) = L_1(\bar{\beta}) + \nabla L_1(\bar{\beta})^T (\beta - \bar{\beta}) + \sum_{t=2}^{\infty} \frac{1}{t!} \nabla^t L_1(\bar{\beta}) (\beta - \bar{\beta})^{\otimes t}$$

Surrogate likelihood 
$$\tilde{L}^1(\beta) = L_1(\beta) + \{\nabla L(\bar{\beta}) - \nabla L_1(\bar{\beta})\}^T \beta$$
 
$$\nabla L(\bar{\beta}) = \frac{1}{\kappa} \sum \nabla L_j(\bar{\beta}); \quad \tilde{\beta} = \operatorname{argmax}_{\beta} \tilde{L}(\beta)$$



# **Simulation Study**

#### Number of sites (K)



#### Real-world Data Application

Investigate the relationship between **COVID-19 viral** (SARS-CoV-2 polymerase chain reaction [PCR] or antigen) **test positivity** and the **symptoms and conditions associated with Long-COVID** in **children** 

- Use case 1: Centralized data from PEDSnet
  - Nine children's hospitals across the nation
  - Sample size: 184,501



- Use case 2: Decentralized datasets
  - 12 sites: PEDSnet (9), OHDSI (2), OneFlorida (3)
  - Sample size: 452,160

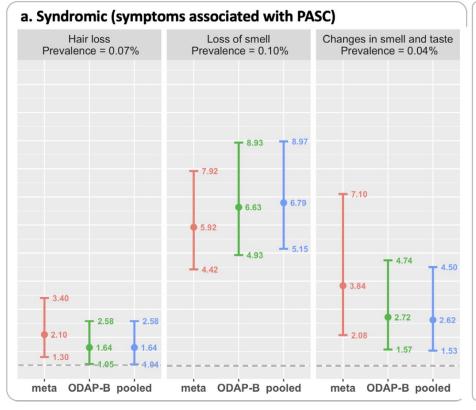


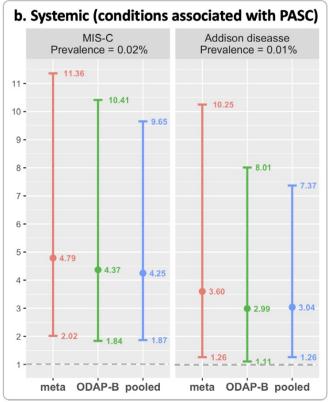




#### Use Case 1 Result

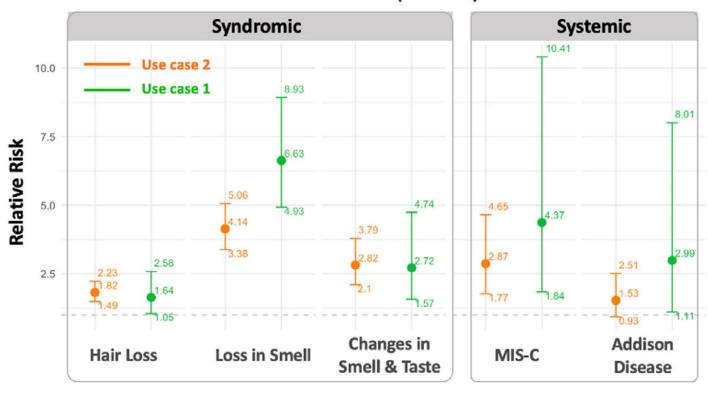
# Results for risk factors: Viral-test Positivity Relative Risk (95% CI)





# Use Case 1 & Use Case 2 Results Comparison

#### Results for risk factors: Viral-test Positivity Relative Risk (95% CI)



#### **Summary**

- ODAP-B is an effective federated learning algorithm for Poisson regression to study rare binary outcome.
- ODAP-B provides inference on adjusted relative risk with robust variance estimator.
- ODAP-B is easy to implement and applicable to analyze multi-site data

R

PDA R Package: 13300+ downloads since 2020



PDA Github Page: https://github.com/Penncil/pda



Privacy-preserving Distributed Algorithm

PDA website: https://pdamethods.org/



PDA-OTA: https://pda-ota.pdamethods.org/





Statistics in Medicine

RESEARCH ARTICLE OPEN ACCESS

Advancing Interpretable Regression Analysis for Binary Data: A Novel Distributed Algorithm Approach

