

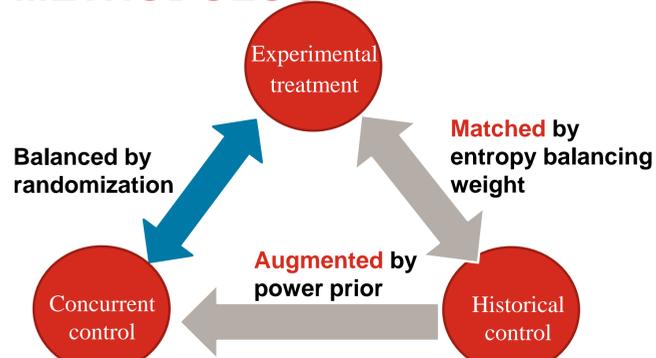
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## INTRODUCTION & OBJECTIVE

- Disproportionate randomized-controlled trials (RCTs) are more appealing and feasible than adequate RCTs for:
  - Pediatric populations, orphan medicines, ethical issues or concerns
- Augmenting the disproportionate RCTs with the real world data (RWD) is prevalent
- OBJECTIVE: Augment the disproportionate randomized-controlled trial by leveraging the real world data (RWD).**
  - Adjust the heterogeneity of the data from the current trial and historical studies
  - Control the amount of historical information borrowing

## METHODOLOGY



**Key Assumptions:**  
Patients in the control arm in current RCT and historical studies received the same active control treatment or placebo.

## SIMULATION SETUP

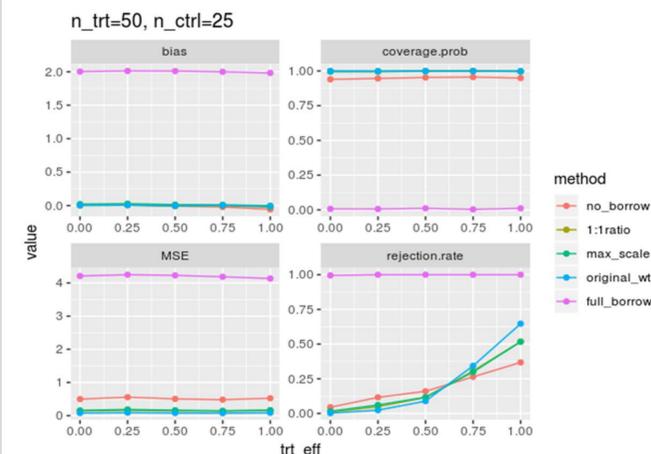
### Data generation:

- Generate two **heterogeneous** patients populations  $X_{ki}$
- Sample from these two populations to form current RCT and historical trial patients.
- Generate the outcome data  $Y_{ki} \sim N(\tilde{\beta}^T X_{ki} + \beta_1 * Z_{ki}, \sigma^2)$  with  $\tilde{\beta}$  as the prognostic effects.

### Operating setting:

- For current RCT,  $N_{trt} : N_c = 2:1$ ;
- For historical control patients,  $N_{c0} : N_c = 3:1$ ;
- Consider treatment effect  $\beta_1 = 0, 0.25, 0.5, 0.75, 1$ ;
- Define the decision rule as  $\Pr(\beta_1 > 0 | D_1, D_0) > 0.95$

## KEY RESULT



## CONCLUSIONS

- Poor estimation when borrowing historical information without adjusting the heterogeneity.
- The proposed method improves the efficiency of estimation (lower MSE and higher coverage probability), especially when the current study is a small scaled trial.
- The proposed method results in a low type 1 error and an increasing power when effect size is larger.
- The amount of information borrowing is flexible and can be calibrated by a scalar.
- Enables a more comprehensive evaluation on treatment effect than frequentist inference approach.

## Notations

### Patient-level data

$$D_{ki} = (Y_{ki}, Z_{ki}, X_{ki}), k = 0, 1; i = 1, \dots, N_k$$

$Y_{ki}$  = Outcome variable;

$Z_{ki}$  = Treatment group indicator: 1-treatment, 0- control

$X_{ki}$  = baseline characteristics covariates vector which may confound the outcome  $Y_{ki}$

$k$  = data source indicator: 0- historical data; 1-current RCT data

## Entropy Balancing Weight ( $w_{ki}$ )

$$\min_{w_{ki}} H(w) = \sum_{\{i|k=0\}} h(w_{ki}), \quad h(\cdot) \text{ is the distance metric}$$

Subject to balance and normalization constraints:

$$\sum_{\{i|k=0\}} w_{ki} c_{ri}(X_{ki}) = m_r \quad \text{with } r \in 1, \dots, R \text{ and } m_r \text{ as a set of balance constraints}$$

and

$$\sum_{\{i|k=0\}} w_{ik} = c_0 \quad w_{ki} \geq 0 \text{ for all } i$$

Compared to propensity score based method, entropy balancing is **advantageous** in terms of

- No model is required.
- No covariates balance check is needed.
- Achieves a better covariate distributions balance.

## Likelihood

**Outcome variables:**  $Y_{ki} \sim N(\mu_{ki}, \sigma^2)$ ,

where  $\mu_{ki} = \beta_0 + \beta_1 * Z_{ki}$  (after the confounding covariates vector  $X_{ki}$  is **balanced** between the treatment and control patients)

- $\beta_0$  – control mean;  $\beta_1$  – treatment effect
- Non-informative priors on  $\beta_1$  and  $\sigma^2$
- Power prior on  $\beta_0$

## Power Prior

$$\pi(\beta_0 | \alpha_0, D_0) \propto \prod_{i=1}^{N_0} L(\beta_0 | D_{0,i})^{\alpha_{0i}} \pi(\beta_0)$$

**The discounting factor**  $\alpha_0 = (w_{0,1}, \dots, w_{0,N_0}) \times c$

Entropy balancing weight      Scalar to adjust the amount of information borrowing

Consider the following **weight scaling**:

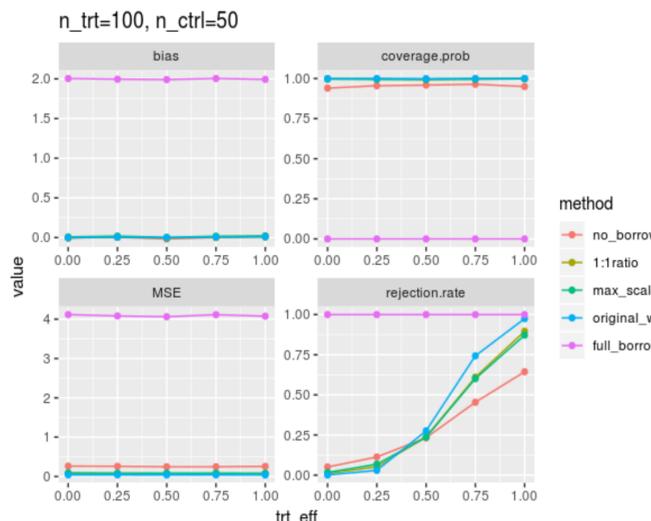
- $c = 1$ , no scaling
- $c = 1/\max(w_{0,1}, \dots, w_{0,N_0})$ , scaling the weight to be between 0 and 1.
- $c = \tilde{c}_0$ , such that 1:1 trt vs. augmented control allocation

## Joint posterior distribution

$$q(\beta, \sigma^2 | D_1, D_0, \alpha_0) \propto L(\beta, \sigma^2 | \tilde{X}, Y) \pi(\beta, \sigma^2) \left[ \prod_{i=1}^{N_0} f(Y_{0,i} | \beta_0, \sigma^2)^{\alpha_{0i}} \right]$$

- $\beta = (\beta_0, \beta_1)$
- $\tilde{X} = (\tilde{X}_1, \dots, \tilde{X}_{N_1})'$  with  $\tilde{X}_i = (1, Z_i)'$  where  $N_1$  is the sample size of the current RCT.
- $D_1, D_0$  denote the data from current RCT and RWD from historical studies, respectively.
- $Y_{0,i}$  is the outcome of  $i$ -th patient from the historical control.

## MORE RESULT



**Note:** R package “**ebal**” was used to calculate the entropy balancing weights in the simulation.

## REFERENCES

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**Disclosures:** GY, YB, and MG are full employees of Eli Lilly and Company.