

Statistical Inference for Maximin Effects: Identifying Stable Associations across Multiple Studies

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Overview of talk

1 Group Distributionally Robust Models

2 Identification and Inference Challenge

3 Method and Theory

4 Numerical Studies

Multi-source data is everywhere

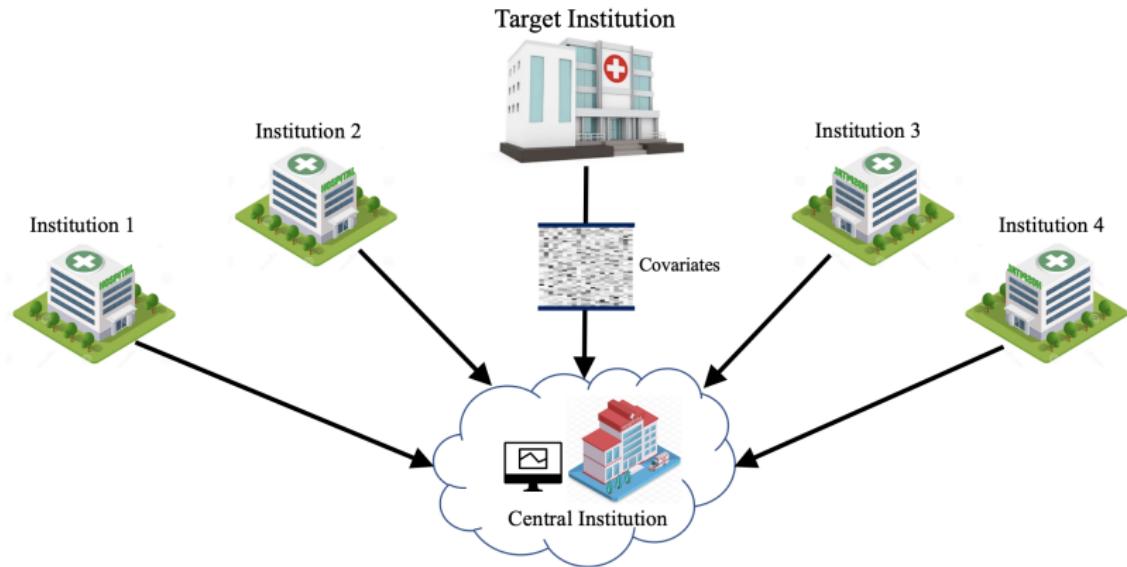


Figure: Figure credited to the Cai Lab at Harvard.

Invariance from Multi-source Data

- Data heterogeneity: sub-populations, healthcare centers and environments.
- Standard model averaging: data merging, meta-analysis, and federated learning
- Today's focus: group distributionally robust models
 - Heterogeneity: allowing for distributional shifts
 - Generalizability: capturing stable associations shared across different sites

Setups

For $1 \leq l \leq L$, the training data $\{X_i^{(l)}, Y_i^{(l)}\}_{1 \leq i \leq n_l}$ follows,

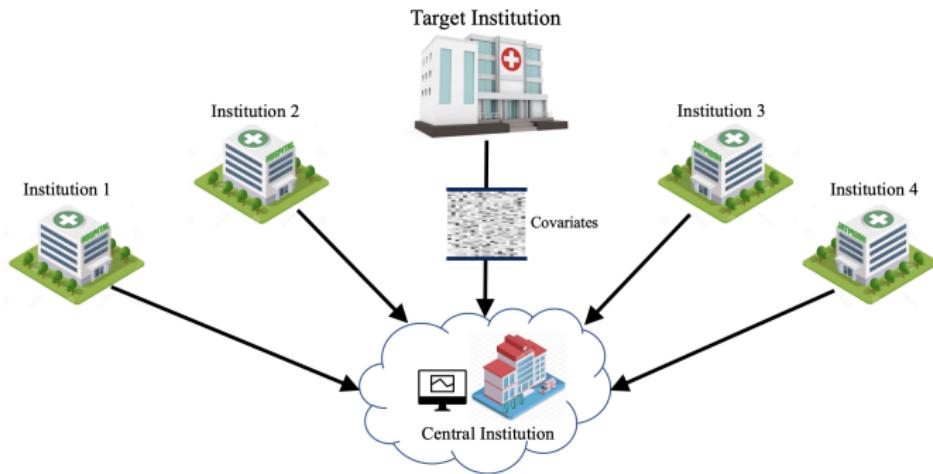
$$X_i^{(l)} \stackrel{\text{i.i.d.}}{\sim} \mathbb{P}_X^{(l)}, \quad Y_i^{(l)} | X_i^{(l)} \stackrel{\text{i.i.d.}}{\sim} \mathbb{P}_{Y|X}^{(l)} \quad \text{for } 1 \leq i \leq n_l.$$

We consider the target population,

$$X_i^Q \stackrel{\text{i.i.d.}}{\sim} \mathbb{Q}_X, \quad Y_i^Q | X_i^Q \stackrel{\text{i.i.d.}}{\sim} \mathbb{Q}_{Y|X} \quad \text{for } 1 \leq i \leq N_Q,$$

- ① No observations of $\{Y_i^Q\}_{1 \leq i \leq N_Q}$.
- ② Covariate shift: \mathbb{Q}_X is different from $\{\mathbb{P}_X^{(l)}\}_{1 \leq l \leq L}$.
- ③ $\mathbb{Q}_{Y|X}$ is different from $\{\mathbb{P}_{Y|X}^{(l)}\}_{1 \leq l \leq L}$.

A class $\mathcal{C}(\mathbb{Q}_X)$ of target populations



$$\mathcal{C}(\mathbb{Q}_X) = \left\{ \mathbb{T} = (\mathbb{Q}_X, \mathbb{T}_{Y|X}) : \mathbb{T}_{Y|X} = \sum_{l=1}^L q_l \cdot \mathbb{P}_{Y|X}^{(l)} \text{ with } q \in \Delta_L \right\},$$

where $\Delta_L := \left\{ q \succ 0 : \sum_{l=1}^L q_l = 1 \right\}$.

Group Distributionally Robust Models

If the test data $\{X_i, Y_i\} \sim \mathbb{T}$, define the reward function for β ,

$$\mathbf{E}_{\mathbb{T}} Y_i^2 - \mathbf{E}_{\mathbb{T}}(Y_i - X_i^\top \beta)^2.$$

The worst-case or adversarial reward,

$$R_{\mathbb{Q}}(\beta) = \min_{\mathbb{T} \in \mathcal{C}(\mathbb{Q}_X)} \left\{ \mathbf{E}_{\mathbb{T}} Y_i^2 - \mathbf{E}_{\mathbb{T}}(Y_i - X_i^\top \beta)^2 \right\}.$$

The group distributionally robust model

$$\beta^*(\mathbb{Q}) := \arg \max_{\beta \in \mathbb{R}^p} R_{\mathbb{Q}}(\beta).$$

Sagawa, S., Koh, P. W., Hashimoto, T. B., & Liang, P. (2019). Distributionally robust neural networks for group shifts: On the importance of regularization for worst-case generalization.

Covariate-shift Maximin Effect

If $\mathbb{P}_X^{(l)} = \mathbb{Q}_X$, then $\beta^*(\mathbb{Q})$ is reduced to the maximin effect

$$\beta^* := \arg \max_{\beta \in \mathbb{R}^p} R(\beta) \quad (1)$$

$$R(\beta) = \min_{1 \leq l \leq L} \left\{ \mathbf{E}[Y_1^{(l)}]^2 - \mathbf{E}[Y_1^{(l)} - (X_1^{(l)})^\top \beta]^2 \right\}.$$

Meinshausen, N., & Bühlmann, P. (2015). Maximin effects in inhomogeneous large-scale data. The Annals of Statistics, 43(4), 1801-1830.

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Identification

$$Y_{n_l \times 1}^{(l)} = X_{n_l \times p}^{(l)} b_{p \times 1}^{(l)} + \epsilon_{n_l \times 1}^{(l)}, \quad \text{for } 1 \leq l \leq L.$$

Theorem (Meinshausen & Bühlmann (2015), G. (2020))

If $\lambda_{\min}(\Sigma^{\mathbb{Q}}) > 0$, then $\beta^*(\mathbb{Q})$ is identified as

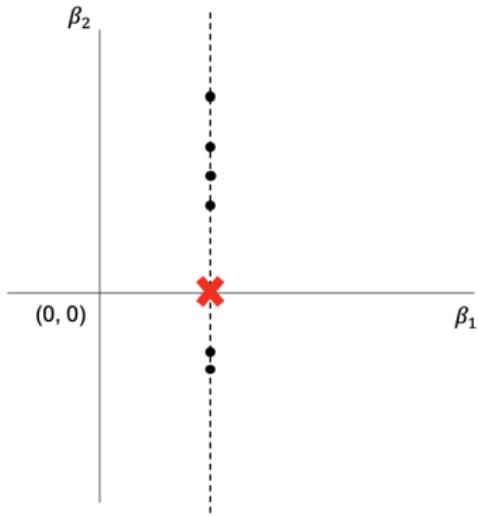
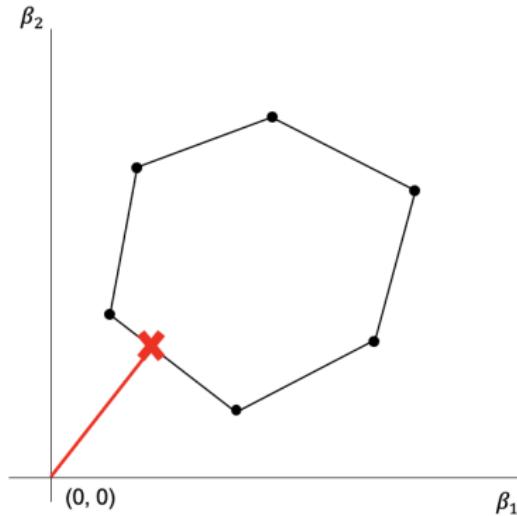
$$\beta^*(\mathbb{Q}) = \sum_{l=1}^L [\gamma^*(\mathbb{Q})]_l b^{(l)} \quad \text{with} \quad \gamma^*(\mathbb{Q}) = \arg \min_{\gamma \in \Delta^L} \gamma^T \Gamma^{\mathbb{Q}} \gamma \quad (2)$$

where $\Delta^L = \{\gamma \in \mathbb{R}^L : \gamma_j \geq 0, \sum_{j=1}^L \gamma_j = 1\}$ is the simplex over \mathbb{R}^L and

$$\Gamma_{lk}^{\mathbb{Q}} = (b^{(l)})^T \Sigma^{\mathbb{Q}} b^{(k)} \quad \text{for } 1 \leq l, k \leq L,$$

with $\Sigma^{\mathbb{Q}} = \mathbf{E} X_i^{\mathbb{Q}} [X_i^{\mathbb{Q}}]^T$.

Generalizability: stable associations



Meinshausen & Bühlmann (2015)

Asymptotic normality fails

For two groups, the optimal aggregation weight $(\gamma_1^*, 1 - \gamma_1^*)^\top$ with

$$\gamma_1^* = \min \left\{ \max \left\{ \frac{\Gamma_{22}^Q - \Gamma_{12}^Q}{\Gamma_{11}^Q + \Gamma_{22}^Q - 2\Gamma_{12}^Q}, 0 \right\}, 1 \right\}.$$

$$\hat{\gamma}_1 = \min \left\{ \max \left\{ \frac{\hat{\Gamma}_{22}^Q - \hat{\Gamma}_{12}^Q}{\hat{\Gamma}_{11}^Q + \hat{\Gamma}_{22}^Q - 2\hat{\Gamma}_{12}^Q}, 0 \right\}, 1 \right\}.$$

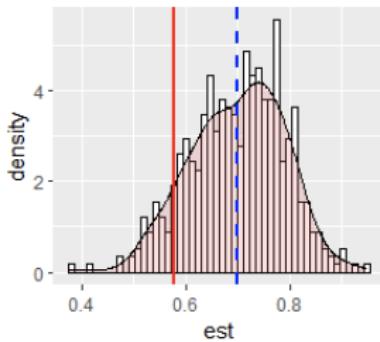
① Non-regularity:

$$\frac{\Gamma_{22}^Q - \Gamma_{12}^Q}{\Gamma_{11}^Q + \Gamma_{22}^Q - 2\Gamma_{12}^Q} \approx 0 \quad \text{or} \quad 1.$$

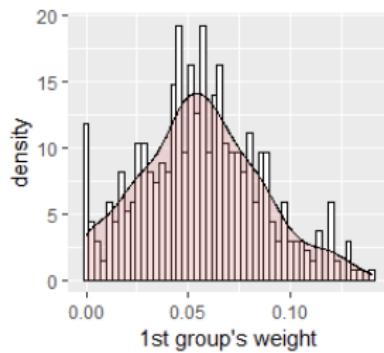
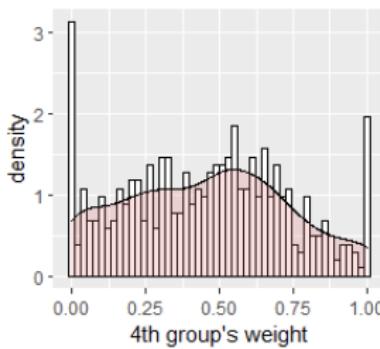
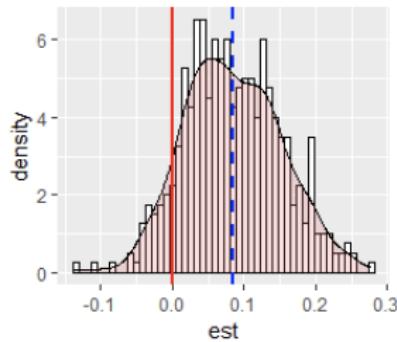
② Instability:

$$\Gamma_{11}^Q + \Gamma_{22}^Q - 2\Gamma_{12}^Q \approx 0.$$

Non-regularity + Instability



Non-regularity



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Two Sources of Randomness

Target parameter: $x_{\text{new}}^T \beta^* = \sum_{l=1}^L \gamma_l^* \cdot x_{\text{new}}^T b^{(l)}$.

Plugin estimator: $\widehat{x}_{\text{new}}^T \beta^* = \sum_{l=1}^L \widehat{\gamma}_l \cdot \widehat{x}_{\text{new}}^T b^{(l)}$

- ① $\widehat{x}_{\text{new}}^T b^{(l)} - x_{\text{new}}^T b^{(l)}$ is asymptotically normal
- ② Even if $\widehat{\Gamma}^Q - \Gamma^Q$ is asymptotically normal, $\widehat{\gamma}_l$ may be non-normal

$$\widehat{\gamma} = \arg \min_{\gamma \in \Delta^L} \gamma^T \widehat{\Gamma}^Q \gamma$$

ReSampling

Note that

$$\text{vecl}(\widehat{\Gamma}^{\mathbb{Q}}) - \text{vecl}(\Gamma^{\mathbb{Q}}) \stackrel{d}{\approx} \mathcal{N}(\mathbf{0}, \widehat{\mathbf{V}}).$$

We resample

$$\left\{ \text{vecl}(\widehat{\Gamma}^{[m]}) \right\}_{1 \leq m \leq M} \stackrel{i.i.d.}{\sim} N \left(\text{vecl}(\widehat{\Gamma}^{\mathbb{Q}}), \widehat{\mathbf{V}} + d_0/n \cdot \mathbf{I} \right),$$

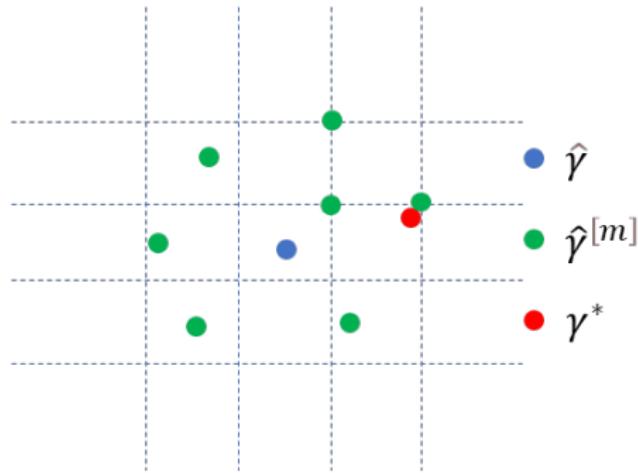
with $d_0 = \max \left\{ \tau \cdot n \|\widehat{\mathbf{V}}\|_{\infty}, 1 \right\}.$

Construct the sampled weight vector as

$$\widehat{\gamma}^{[m]} = \arg \min_{\gamma \in \Delta^L} \gamma^T \widehat{\Gamma}_+^{[m]} \gamma.$$

Xie, M. G., & Singh, K. (2013). Confidence distribution, the frequentist distribution estimator of a parameter: A review. International Statistical Review, 81(1), 3-39.

Sampling property



There exists $1 \leq m^* \leq M$ such that

$$\hat{\gamma}^{[m^*]} \approx \gamma^*.$$

Sampling property

$$\widehat{x_{\text{new}}^T \beta}^{[m]} = \sum_{l=1}^L \widehat{\gamma}_l^{[m]} \cdot \widehat{x_{\text{new}}^T b^{(l)}} \quad \text{for } 1 \leq m \leq M,$$

$$\widehat{x_{\text{new}}^T \beta}^{[m^*]} - x_{\text{new}}^T \beta = \sum_{l=1}^L (\widehat{\gamma}_l^{[m^*]} - \gamma_l^*) \cdot \widehat{x_{\text{new}}^T b^{(l)}} + \sum_{l=1}^L \gamma_l^* \cdot (\widehat{x_{\text{new}}^T b^{(l)}} - x_{\text{new}}^T b^{(l)}).$$

With $m = m^*$,

- The uncertainty of $\sum_{l=1}^L (\widehat{\gamma}_l^{[m^*]} - \gamma_l^*) \cdot \widehat{x_{\text{new}}^T b^{(l)}}$ is negligible
- Quantify the uncertainty of $\sum_{l=1}^L \gamma_l^* \cdot (\widehat{x_{\text{new}}^T b^{(l)}} - x_{\text{new}}^T b^{(l)})$

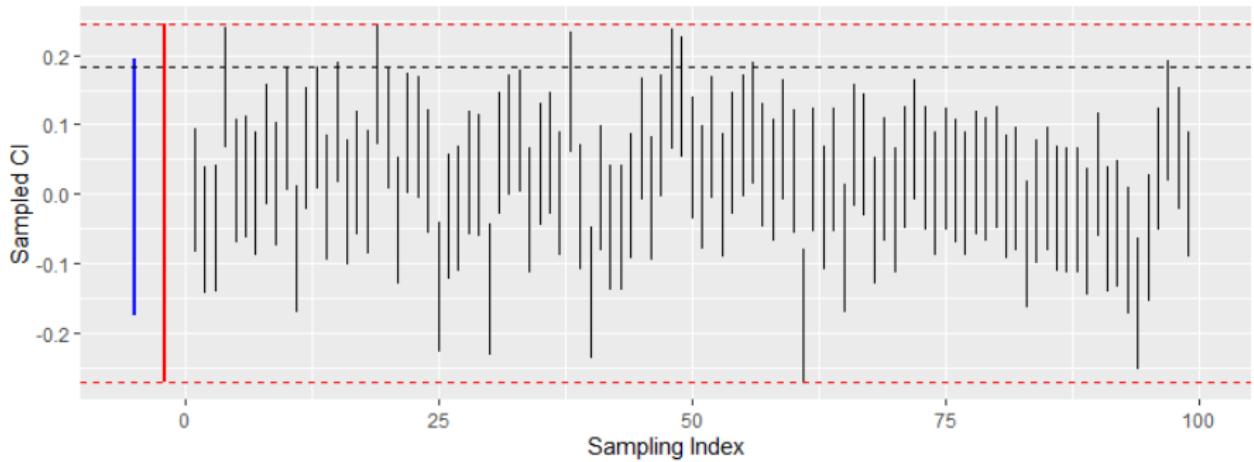
Proposed CI

For $1 \leq m \leq M$,

$$\text{Int}_{\alpha}^{[m]}(x_{\text{new}}) = \left(\widehat{x_{\text{new}}^{\top} \beta}^{[m]} - z_{1-\alpha/2} \widehat{s\epsilon}^{[m]}(x_{\text{new}}), \widehat{x_{\text{new}}^{\top} \beta}^{[m]} + z_{1-\alpha/2} \widehat{s\epsilon}^{[m]}(x_{\text{new}}) \right)$$

$$\widehat{s\epsilon}^{[m]}(x_{\text{new}}) = 1.01 \sqrt{\sum_{l=1}^L [\widehat{\gamma}_{\delta}^{[m]}]_l^2 \frac{\widehat{\sigma}_l^2}{n_l^2} [\widehat{V}^{(l)}]^{\top} (X^{(l)})^{\top} X^{(l)} \widehat{V}^{(l)}}.$$

$$\text{CI}_{\alpha}(x_{\text{new}}^{\top} \beta^*) = \cup_m \text{Int}_{\alpha}^{[m]}(x_{\text{new}}).$$



- red: our proposal.
- blue: asymptotic normal CI with the oracle SE.
- **The confidence interval is of a parametric length.**

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Normality CI

We introduce the normality CI as

$$(\widehat{x}_{\text{new}}^T \widehat{\beta}^* - 1.96 \cdot \widehat{SE}, \widehat{x}_{\text{new}}^T \widehat{\beta}^* + 1.96 \cdot \widehat{SE}), \quad (3)$$

where \widehat{SE} denotes the sample SE calculated based on 500 simulations.

Setting	$\mathbb{I}(\delta)$	Coverage		Length		Length Ratio
		normality	Proposed	normality	Proposed	
(I-1)	3.368	0.700	0.960	0.352	0.597	1.693
(I-2)	3.707	0.818	0.978	0.320	0.543	1.699
(I-3)	3.182	0.748	0.970	0.352	0.588	1.673
(I-4)	1.732	0.770	0.956	0.520	0.796	1.532
(I-5)	1.857	0.796	0.978	0.445	0.710	1.594
(I-6)	1.987	0.710	0.980	0.480	0.832	1.732
(I-7)	0.029	0.848	0.985	0.250	0.507	2.028
(I-8)	0.031	0.758	0.981	0.262	0.530	2.020
(I-9)	0.010	0.830	0.988	0.690	1.264	1.832
(I-10)	0.030	0.940	0.988	0.232	0.315	1.354

Bootstrap or Sub-sampling

Setting $p = 30$	normality	m-out-of-n subsampling				m-out-of-n bootstrap					Proposed	
		$m = 200$	$m = 300$	$m = 400$	$m = 500$	$m = 200$	$m = 300$	$m = 400$	$m = 500$	$m = 1000$	Cov	L-ratio
(I-1)	0.686	0.380	0.390	0.380	0.408	0.432	0.460	0.502	0.490	0.536	0.956	1.678
(I-2)	0.808	0.418	0.456	0.474	0.446	0.456	0.510	0.532	0.546	0.602	0.990	1.723
(I-3)	0.770	0.392	0.494	0.464	0.440	0.482	0.480	0.516	0.544	0.634	0.984	1.766
(I-4)	0.816	0.620	0.668	0.670	0.668	0.672	0.694	0.710	0.700	0.794	0.990	1.686
(I-5)	0.790	0.612	0.626	0.626	0.594	0.628	0.664	0.702	0.732	0.732	1.000	1.843
(I-6)	0.806	0.590	0.636	0.654	0.632	0.626	0.682	0.698	0.712	0.760	0.994	1.833
(I-7)	0.824	0.914	0.932	0.908	0.888	0.890	0.950	0.934	0.950	0.952	0.996	5.078
(I-8)	0.888	0.912	0.934	0.884	0.856	0.914	0.916	0.932	0.932	0.958	0.996	4.017
(I-9)	0.900	0.834	0.822	0.788	0.778	0.914	0.914	0.916	0.862	0.914	0.996	2.061
(I-10)	0.954	0.956	0.904	0.866	0.836	0.956	0.948	0.962	0.964	0.942	1.000	1.725

Subsampling/bootstrap do not solve non-regular inference problems.

Andrews, D. W. (2000). Inconsistency of the bootstrap when a parameter is on the boundary of the parameter space. *Econometrica*, 399–405.

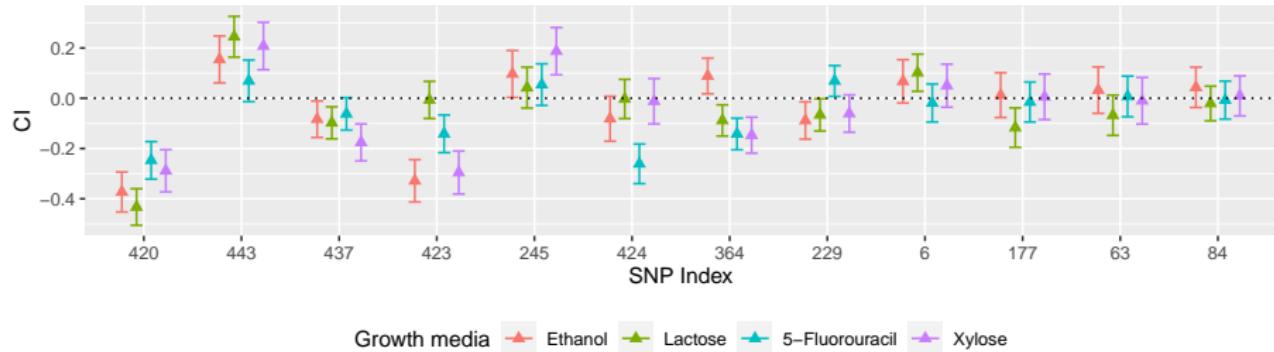
Applications to GWAS

- Study the yeast colony growth under different growth media.
- Outcome: end-point colony sizes under different growth media.
- $p = 513$ genetic markers
- The study is based on $n = 1,008$ yeast segregants.
- Training media: "Ethanol", "Lactose", "5-Fluorouracil", "Xylose"

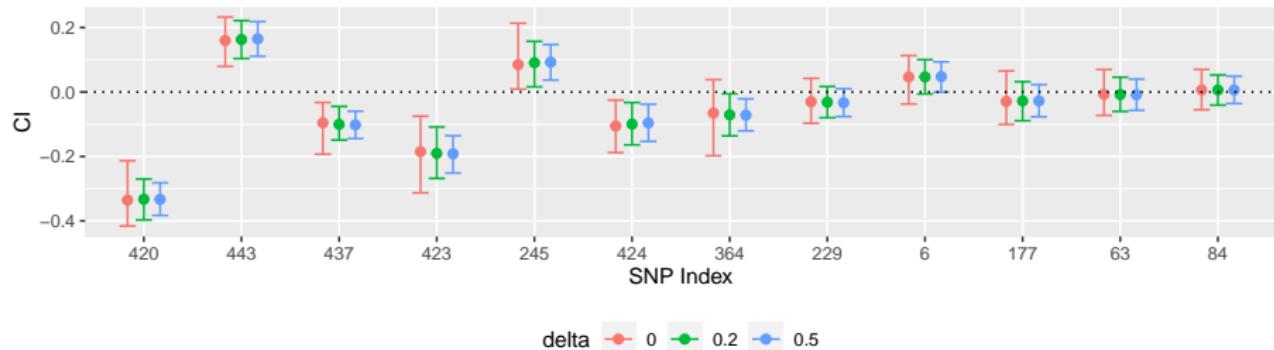
Bloom, J. S., I. M. Ehrenreich, W. T. Loo, T.-L. V. Lite, and L. Kruglyak (2013). Finding the sources of missing heritability in a yeast cross. *Nature* 494 (7436), 234–237.

Stable associations: causal?

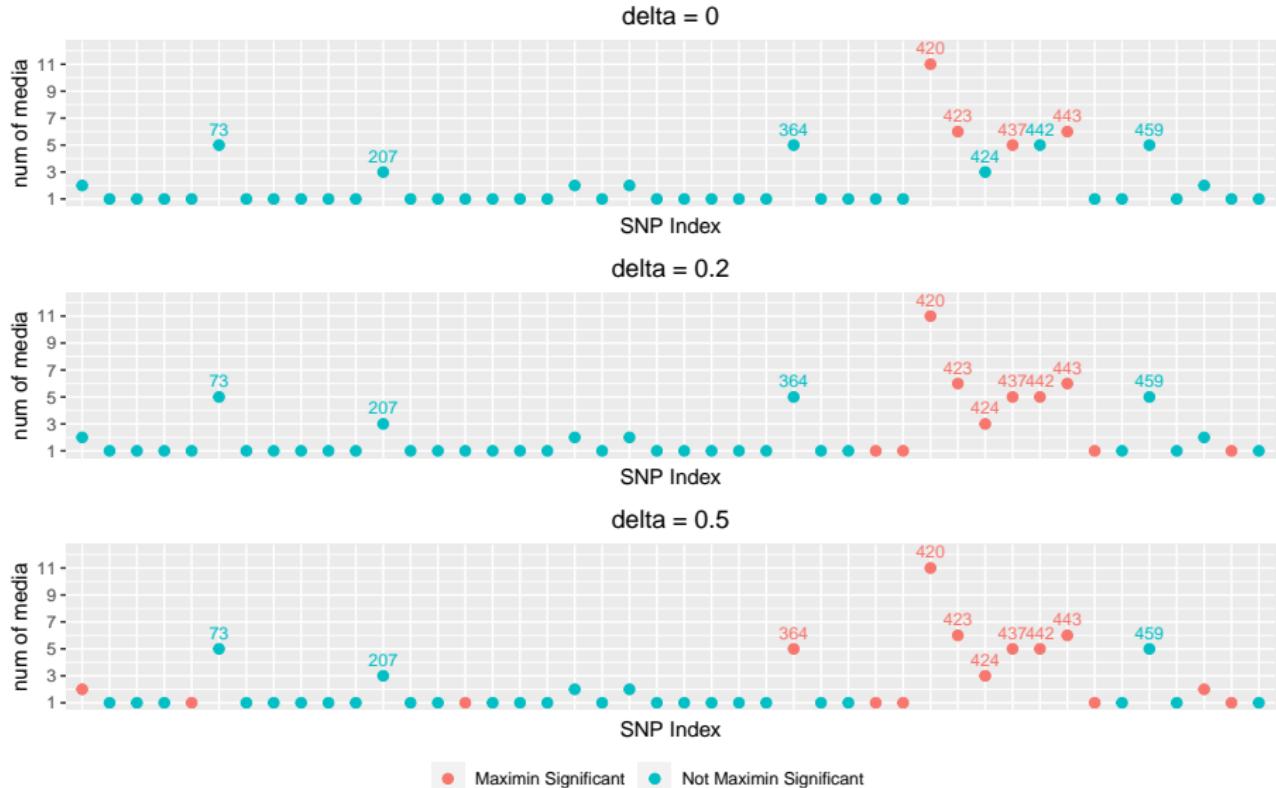
CIs for the regression coefficients for different growth media



CIs for the maximin effect



Generalizability?



Surv-Maximin

- Study the COVID-19 mortality risk prediction.
- Analyze 15 participating health care centers across three countries (France, Germany, and the U.S.).
- The proposed SurvMaximin algorithm achieves higher accuracy than the existing methods.
- Privacy-preserving.

Wang, X., Zhou, H., . . . , 4CE, Avillach, P., **Guo, Z.**, & Cai, Tianxi. (2022) SurvMaximin: Robust Federated Approach to Transporting Survival Risk Prediction Models. *Journal of biomedical informatics* 134 (2022): 104176.

Take home message

Multi-source data

- heterogeneity
- generalizability
- privacy-preserving.
- Non-standard inference: sampling
- ...

Generalizable model

- Maximin effect
- Group distributionally robust model
- ...

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Guo, Z. (2020). Statistical Inference for Maximin Effects: Identifying Stable Associations across Multiple Studies. *arXiv preprint arXiv:2011.07568*.

R package MaximinInfer

Thank you!