# Statistical Inference with Screening and Selection

Isaiah Andrews Tuesday, May 23

## **Screening and Selection**

- Frequentist statistical guarantees control performance under repeated sampling
  - That is, if we could draw the data multiple times in a given situation, certain properties would hold on average across data realizations
- The way in which statistical procedures are commonly applied often doesn't match the sampling thought experiment
  - · We might only write up/publish certain findings
  - We might choose the target for inference based on the data
- In such cases, standard statistical procedures can yield non-standard behavior

- To illustrate, imagine that we conduct a randomized trial of a job training program
- Usual statistical procedure:
  - Compare average outcomes in treatment and control groups
  - Conclude that treatment has an effect if the difference in average outcomes is large relative to the standard error
- Analogously, we can analyze the effect of treatment within observable subgroups of the sample, e.g. based on location or prior employment history
- Usual statistical guarantee: under regularity conditions, if treatment in fact has no effect, we will mistakenly conclude that there is an effect at most α (e.g. 5%) of the time

## **Screening and Selection**

- The usual statistical guarantee fixes a procedure (e.g. run the experiment and take the difference-in-means) and asks how it performs over repeated draws of the data
- Empirical practice often differs from this idealized description
  - Our experimental result may only be written up or published if it is positive and statistically significant ⇒ screening/publication bias
  - We might focus our analysis on the subgroup with the largest effect
     ⇒ selection bias
- There is an active literature in statistics and related fields which aims to correct for these issues
  - My goal today: provide a brief review of this literature, some of the tools it suggests, and some of the questions that it raises

# Decision Theory Without Screening or Selection

## **Decision Theory Without Screening or Selection**

- Suppose we observe  $X \sim F(\mu)$  for an unknown parameter  $\mu$
- For today, will specialize to  $X \sim N(\mu, \Sigma)$  for  $\mu \in \mathbb{R}^J$  and  $\Sigma$  known
  - For X a vector of estimates based on underlying, potentially non-normal observations, justified in many contexts by the central limit theorem
- Suppose we are interested in a parameter  $\theta = \mathbf{v}' \mu \in \mathbb{R}$
- For a set of possible actions A, and a loss function L, we want to choose a decision rule δ (X) to achieve a low expected loss

 $E_{\mu}\left[L\left(\delta\left(X\right),\theta\right)\right].$ 

We may also require this rule to satisfy some additional constraints

- In the job-training experiment, the vector X could collect treatment-control differences across J demographic subgroups
  - So long as the number of trial participants in each subgroup is large, the central limit theorem justifies the approximation

$$X \sim N(\mu, \Sigma)$$

for  $\Sigma$  a diagonal matrix

- We can consider different target parameters  $\theta$  in this context
  - For  $\omega_j$  the population share of group *j* and  $v = (\omega_1, ..., \omega_J)'$ ,

$$\theta = \mathbf{v}' \mu = \sum_j \omega_j \mu_j$$

captures the average treatment effect in the population

• For  $v = e_j = (0, .., 0, 1, 0, ..., 0)$  the *j*th standard basis vector,

$$\theta = \mathbf{v}' \mu = \mu_j$$

captures the average treatment effect in subgroup *j* 

## Loss Functions and Constraints

• For estimation we can take  $A = \mathbb{R}$  and consider squared-error loss  $L(a, \theta) = (a - \theta)^2$ , so

$$E_{\mu}\left[L\left(\delta\left(X\right),\theta\right)\right]=E_{\mu}\left[\left(\delta\left(X\right)-\theta\right)^{2}\right]$$

corresponds to mean squared error

We may further impose unbiasedness or median-unbiasedness,

$$E_{\mu}[\delta(X)] = \theta \text{ or } Med_{\mu}(\delta(X) > \theta) = \frac{1}{2} \text{ for all } \mu$$

 To quantify uncertainty we might focus on confidence intervals, taking A to be the set of closed intervals in ℝ, define L (a, θ) = |a| as the length of a, and impose a coverage constraint,

$$Pr_{\mu} \{ \theta \in \delta(X) \} \geq 1 - \alpha \text{ for all } \mu$$

potentially along with other constraints

These problems have well-known solutions

• The maximum likelihood estimator

$$\hat{\theta} = v' X$$

is the best (median-)unbiased estimator for  $\theta,$  in the sense that for any other (median-)unbiased estimator  $\tilde{\theta},$ 

$$E_{\mu}\left[\left(\hat{ heta}- heta
ight)^{2}
ight]\leq E_{\mu}\left[\left( ilde{ heta}- heta
ight)^{2}
ight]$$
 for all  $\mu$ 

• Similarly, for  $\sigma_{\hat{\theta}} = \sqrt{v' \Sigma v}$  the standard error of  $\hat{\theta}$ , confidence intervals of the form

$$\left[\hat{\theta} \pm \boldsymbol{c} \cdot \boldsymbol{\sigma}_{\hat{\theta}}\right]$$

are optimal in various senses

• c = 1.96 for gives the standard 95% confidence interval.

## **Screening and Selection**

- The decision-theoretic setup above made two assumptions
  - 1. We care about performance on average across all realizations of X
  - 2. The target parameter  $\theta = v'\mu$  is the same for all realizations of *X*

which can fail in practice

- I'll refer to failures of (1) as **screening** and failures of (2) as **selection**
- Warning: useful shorthand for today, but these are not consistently adopted terms in literature

# Screening

## **Screening Problems**

- Above, we averaged performance over all realizations of X
- Sometimes, however, we may only care about performance over a subset of data realizations
- To formalize this, suppose that for a screening variable S
  - 1. The conditional distribution of S|X does not depend on  $\mu$
  - 2. We only care about behavior conditional on S = 1, e.g.

 $E_{\mu}\left[L\left(\delta\left(X\right),\theta\right)|S=1\right], \ E_{\mu}\left[\delta\left(X\right)|S=1\right], \ Pr_{\mu}\left\{\theta\in\delta\left(X\right)|S=1\right\}$ 

- In the randomized trial example, suppose θ corresponds to the average treatment effect over the population
- Randomization ensures that θ̂ = ν'X is unbiased for θ on average across data realizations
  - But not all estimates  $\hat{\theta}$  are equally likely to be published
  - An extensive literature expresses concern about, and provides evidence of, publication bias
  - A few recent examples include Open Science Collaboration (2015), Bruns and Ioannidis (2016), and Camerer et al. (2016)

 To study publication bias in this example, let S = 1 be an indicator for the event that a given estimate θ̂ gets published,

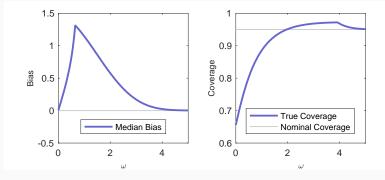
$$oldsymbol{S} = oldsymbol{1} \left\{ \mathsf{Estimate} \; \hat{ heta} \; \mathsf{gets} \; \mathsf{published} 
ight\}$$

- We assumed that that the conditional distribution S|X does not depend on  $\theta$
- Means that publication decisions depend only on estimates, and not on the underlying parameters once we hold the experimental results fixed
- Publication decisions could depend on many factors
  - Preference for positive results
  - Preference for surprising results
  - Preference for results consistent with previous literature

- Here, I'll focus on an example from Andrews and Kasy (2019)
  - Preference for statistically significant results: many papers in the literature point to this
  - Specifically, let Z = θ̂/σ<sub>θ̂</sub> ~ N(ω, 1) denote the estimate, standardized to have variance one
  - Suppose results that are significantly different from zero at the 5% level (i.e |Z| > 1.96) are 10 times more likely to be published than are statistically insignificant results

$$Pr\left\{S=1|X
ight\}\propto egin{cases} 1 & ext{if $X$ implies $|Z|>1.96$} \ 0.1 & ext{otherwise} \end{cases}$$

 What is the effect of such screening on the distribution of published results?



## **Screening Problems**

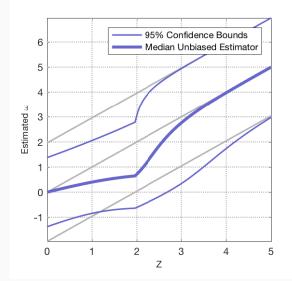
- While I've discussed screening in terms of publication bias, note that it doesn't matter who's doing the screening
  - e.g. authors choosing not to write up results vs. journals choosing not to publish
  - Hence, "screening" as discussed here covers many forms of so-called "p-hacking"
- Moreover, some practices which are recommended on other grounds generate the same issues
  - e.g. dropping models where a specification test suggests the model is incorrect

## **Corrections for Screening**

- Fortunately, some results in statistics provide powerful tools to correct for screening
- Let f<sub>X</sub> (x|µ) denote the density of X without screening. Bayes rule implies that the conditional density of X given S = 1 is

$$f_{X|S=1}\left(x|\mu\right) = \frac{E\left[S|X=x\right]f_{X}\left(x|\mu\right)}{E_{\mu}\left[S\right]}$$

- This implies that if f<sub>X</sub> (x|µ) has exponential family structure (as is true for the normal distribution) then f<sub>X|S=1</sub> (x|µ) does as well
- Results in statistics then deliver optimal median-unbiased estimators, optimal confidence intervals for  $\theta = \mathbf{v}' \mu$



## Selection

## **Selection Problems**

- In screening problems, we only cared about some values of *X*, but always cared about the same target θ
- In selection problems, by contrast, we want to conduct inference on θ<sub>X</sub> = v (X)' μ
  - Hence, we may have a different target for inference for different values of *X*
- Selection problems of this sort have been extensively studied in the recent statistics literature
  - Motivated by model selection concerns: let M<sub>X</sub> be the model selected when realized data are X, and define θ<sub>X</sub> as the target parameter under this model
  - e.g. Berk et al. (2013), Lee et al (2016), Fithian et al. (2017)
- Selection problems also arise outside the context of model selection, however

- Recall that in this example, *X* records the treatment-control differences over *J* different subgroups
- We might be interested in the effect of treatment on the group for whom treatment appears most effective,

$$heta_{X}= extbf{e}_{\hat{j}}^{\prime}\mu,\ \hat{j}=rg\max X_{j}$$

 Alternatively, we might be interested in the average effect of treatment across those subgroups where it appears helpful,

$$\theta_X = v(X)' \mu, \ v_j(X) = \frac{\omega_j \mathbf{1} \{X_j > 0\}}{\sum_j \omega_j \mathbf{1} \{X_j > 0\}}$$

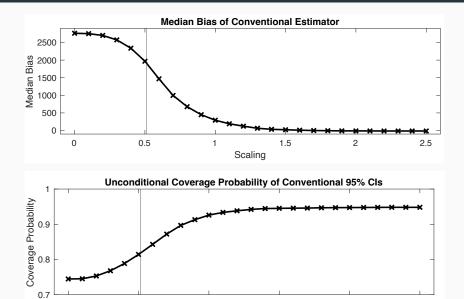
- As with screening, selection generally invalidates conventional inference approaches
- To illustrate, use an example from Andrews et al. (2023)
- Calibrate a simulation based on data from a randomized trial of job-training programs, conducted at 13 different sites
  - JOBSTART Experiment, conducted by US Department of Labor
  - Program at one site appeared most effective
  - This site was subsequently selected as the model for a subsequent replication study conducted at new sites
  - Results in replication turned out to be disappointing

Simulations take the experimental results as starting point

- Scale up/down to vary the heterogeneity in effect sizes across sites
- For a given effect size, simulate draws of experimental results, and ask how selecting the site with the largest estimated effect impacts inference
- Vertical line shows scaling to match unbiased estimate for variance of effects across sites.

0.5

0



Scaling

1

1.5

2

2.5

## **Selection Problems**

- We see that, like screening, selection can invalidate conventional inference procedures
  - Estimated effect sizes are biased upwards
  - · Conventional confidence intervals under-cover
- The selection problem introduces a new wrinkle relative to our analysis so far: the target parameter θ<sub>X</sub> is now random
  - Different target parameters for different data realizations
- This suggests two possible routes forward:
  - We could condition on v (X) to remove this randomness...
  - ... or we could not

## **Conditional Inference**

 By conditioning on v (X), I mean requiring conditional median unbiasedness or conditional coverage

$$\begin{array}{l} \textit{Med}_{\mu}\left(\delta\left(X\right)|v\left(X\right)=\tilde{v}\right)=\tilde{v}'\mu\\ \textit{Pr}_{\mu}\left\{\tilde{v}'\mu\in\delta\left(X\right)|v\left(X\right)=\tilde{v}\right\}=1-\alpha \end{array} \text{ for all } \mu,\tilde{v} \end{array}$$

- However, this immediately returns us to the selection case by defining S = 1 {v (X) = ṽ}
  - Hence, we know how to construct optimal estimators and confidence sets once we condition on the target parameter
- This route was advocated by Fithian et al (2017): Our guiding principle is: The answer must be valid, given that the question was asked.

## **Unconditional Inference**

 Alternatively, we could focus just on unconditional bias and coverage, requiring that

 $\begin{array}{l} \textit{Med}_{\mu}\left(\delta\left(X\right)-v\left(X\right)\right)=0\\ \textit{Pr}_{\mu}\left\{v\left(X\right)'\theta\in\delta\left(X\right)\right\}\geq1-\alpha \end{array} \text{ for all }\mu\end{array}$ 

- Unconditional inference is less demanding
  - Any procedure that is conditionally valid for all v is also unconditionally valid by the law of iterated expectations
  - This also means that the class of unconditionally valid procedures is larger ⇒ may be able to obtain better performance

## **Unconditional Inference**

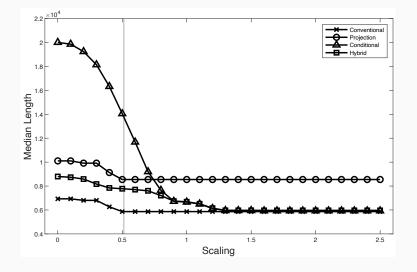
 Berk et al. (2013)'s initial proposal for unconditional inference amounts to forming a joint confidence set for μ, that is, a set CS<sub>μ</sub> = CS<sub>μ</sub>(X) such that

$$Pr_{\mu} \{ \mu \in CS_{\mu} \} \geq 1 - \alpha \text{ for all } \mu,$$

and then forming a confidence set for  $\theta_X$  as

$$\delta\left(\boldsymbol{X}\right) = \left\{\boldsymbol{v}\left(\boldsymbol{X}\right)'\boldsymbol{\mu}:\boldsymbol{\mu}\in\boldsymbol{C}\boldsymbol{S}_{\boldsymbol{\mu}}\right\}$$

- i.e. take the projection of  $CS_{\mu}$  on the dimension of interest
- This ensures (unconditional) coverage, but it can result in confidence sets that are much longer than necessary
- On the other hand, an advantage of this approach is that we don't need to know the function v (·) to implement it suffices to know v (X)
- When *v*(·) is known, Andrews et al. (2023) propose a hybrid approach that combines projection and conditioning



# **Open Questions**

## Can We Relax Information Requirements?

- The available techniques to correct for screening and selection impose substantial information requirements
  - For screening, need to know  $Pr \{S = 1 | X\}$
  - For selection, need to know either v (·) (for conditional and hybrid inference) or the set of possible target parameters θ<sub>X</sub> (for projection inference)

In many contexts, this is too demanding: we do not have an explicit description of what guides our choices

- In some contexts, we may be able to estimate screening or selection rules based on observed choices
  - Andrews and Kasy (2019) do this in the case of publication bias
- In other contexts, we may resort to sample-splitting
  - Screen or select based on part of the data, and use the remainder for inference
- Are there better options?

## How to Think About Screening?

- Screening invalidates conventional inference
  - Motivates suggestions to reduce screening, e.g. pre-analysis plans, registered reports (i.e. pre-result peer review)
- However, this isn't the only option: once the form of screening is known we can correct for it
- Moreover, there are cases where screening seems to be helpful
  - Frankel and Kasy (2022) show that screening in favor of suprising results can be optimal for a journal seeking to inform readers
  - Screening based on specification tests is a common (implicit or explicit) suggestion

When the target parameter is  $\theta_X$ , open questions include:

- Should we condition on the target parameter selected?
- If not, what's the right framework for optimal inference when the target parameter is random?

# Thanks very much!

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