

Debiasing and subsampling/resampling approaches

Patrick Breheny

April 8, 2025

Introduction

- Today's notes will discuss two unrelated approaches to inference:
 - Debiasing, in which we attempt to get around the fact that $\hat{\beta}_j$ is biased by constructing a new statistic $\tilde{\beta}_j$ that is unbiased for β_j
 - Perturbation approaches that use subsampling, resampling, or sample splitting as ways to carry out inference for high-dimensional models
- Both of these are really categories of approaches rather than a specific approach; many ideas have been proposed that fall into each category

Debiasing

- The basic idea behind debiasing is that frequentist inference tends to work well if $\hat{\beta}_j \sim N(\beta_j, \text{SE}^2)$
- Penalized regression estimates obviously do not have this property (with the possible exception of MCP/SCAD), so debiasing approaches construct a new estimate

$$\tilde{\beta}_j = \hat{\beta}_j + \text{adj},$$

for which approximate unbiased normality holds

- Zhang and Zhang (2014)
- van de Geer et al. (2014)
- Javanmard and Montanari (2014)

Implementation

- The adjustment typically takes the form

$$\tilde{\beta} = \hat{\beta} + \hat{\Theta} \frac{1}{n} \mathbf{X}^\top \mathbf{r},$$

where $\hat{\Theta}$ is an estimate of the inverse of $\mathbf{X}^\top \mathbf{X}/n$

- This is easy to understand in the orthogonal case, where $\Theta = \mathbf{I}$ and $\tilde{\beta}$ is simply the OLS estimate
- In high dimensions, however, it is not trivial to estimate Θ and typically involves fitting a new model (e.g., a lasso model) for each feature (treating it as the outcome)

Semi-penalization

- For the sake of this class, let's look at a relatively simpler way to accomplish debiasing: *semi-penalization*
- The idea here is that we can obtain a (more or less) unbiased estimate for β_j by not penalizing it; for example,

$$L(\beta|\mathbf{X}, \mathbf{y}) + \lambda \sum_{k \neq j} |\beta_k|$$

- As far as I know, this idea first appeared in Huang et al. (2013, "SPIDR"); today I'll talk about an approach proposed in Shi et al. (2019), which is very similar in concept but differs in the details

Semi-penalized LRT

- The idea here is actually very similar to the general statistical idea of a likelihood ratio test: we fit constrained and unconstrained models, and then compare their likelihoods
- Specifically, for testing $H_0 : \beta_j = 0$, we would solve for $\hat{\beta}_0$ that minimizes

$$L(\beta_{-j} | \mathbf{X}_{-j}, \mathbf{y}) + \lambda \sum_{k \neq j} |\beta_k|$$

as well as $\hat{\beta}_a$ that minimizes

$$L(\beta | \mathbf{X}, \mathbf{y}) + \lambda \sum_{k \neq j} |\beta_k|$$

Distribution

- It can be shown that (with a number of assumptions), the test statistic

$$2\{\ell(\hat{\beta}_a, \hat{\sigma}^2) - \ell(\hat{\beta}_0, \hat{\sigma}^2)\}$$

follows an approximate χ^2 distribution with 1 degree of freedom, where $\ell(\beta, \sigma^2)$ denotes the log-likelihood

- The error variance can be estimated using any of the methods we have discussed in class, but as in the classical LRT, is based on the unrestricted (alternative) model
- The paper discusses score and Wald tests as well, but we'll only look at the LRT

Remarks

- One of the conditions required to show convergence to the proper distribution is that $\sqrt{np'}(\beta_j^*) \rightarrow 0$ for all $j \in \mathcal{S}$
- This is satisfied for MCP/SCAD, but not the lasso; nevertheless, it seems to me to work reasonably well for the lasso also, so I will go ahead and show those results
- This approach would also seem amenable to constructing confidence intervals, although the article doesn't discuss this
- Another issue is that it would seem reasonable to apply a multiple comparison procedure to the p -values, but this is not discussed in the article, so I'll just present the unadjusted p -values

Results: Example data set (10 largest coefficients)

Feature	Estimate	mfd	SPLRT
A1	0.87	< 0.0001	< 0.0001
A2	-0.77	< 0.0001	< 0.0001
A4	-0.50	< 0.0001	< 0.001
A3	0.42	< 0.0001	< 0.001
A6	-0.35	< 0.001	0.01
A5	0.31	< 0.01	0.54
N39	-0.20	0.33	0.03
N25	0.17	0.48	0.07
N22	0.13	0.78	0.17
B9	0.13	0.75	0.03

Comments

- Results seem more or less similar for the noise variables and most of the “A” variables
- However, B9 and A5 illustrate the key difference:
 - We have convincing evidence that one of them is important according to the marginal approach, which isn't concerned about the possibility of indirect associations
 - This is a major concern for conditional approaches, however – neither variable shows up as significant in the semi-penalized LRT

High-dimensional example: TCGA

- Like several conditional approaches, the semi-penalized LRT works nicely in many low- to medium-dimensional situations, but dramatically loses power in high-dimensional data
- For example, in applying the test to our TCGA data, no genes could be identified as significant: the minimum p -value was 0.14 even without any adjustments for multiple comparisons
- In contrast, 95 features are selected via cross-validation, and 16 of those have a local mfdR under 10%

Sample splitting: Idea

- The rest of today's lecture will focus on using subsampling, resampling, and sample splitting as ways to carry out inference for high-dimensional models
- We begin with the simplest idea: sample splitting

Sample splitting: Idea (cont'd)

Sample splitting involves two basic steps:

- Take half of the data and fit a penalized regression model (e.g., the lasso); typically this involves cross-validation as well for the purposes of selecting λ
- Use the remaining half to fit an ordinary least squares model using only the variables that were selected in step (1)

Sample splitting: Example (step 1)

- Let's split the example data set into two halves, D_1 and D_2 , each with $n = 50$ observations
- Fitting a lasso model to D_1 ($n = 50, p = 60$) and using cross-validation to select λ , we select 16 variables:
 - 6 from category A
 - 1 from category B
 - 9 from category N

Sample splitting: Example (step 2)

- Fitting an ordinary linear regression model to the selected variables ($n = 50$, $p = 16$):
 - 5 “A” features are significant in the $p < 0.05$ sense
 - 0 “B” features were significant
 - 0 “N” features were significant
- We can obtain confidence intervals as well, although note that we only obtain confidence intervals for coefficients selected in step (1)

Sample splitting: Advantages and disadvantages

- The main advantage of the sample splitting approach is that it is clearly valid: all inference is derived from classical linear model theory
- The main disadvantages are:
 - Lack of power due to splitting the sample size in half
 - Potential increase in type I error if important variables are missed in the first stage
 - Results can vary considerably depending on the split chosen

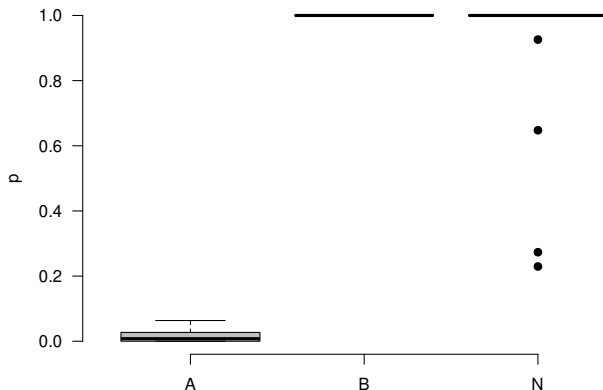
Multiple splits

- An obvious remedy for this final disadvantage is to apply the sample splitting procedure many times and average over the splits
- To some extent, this will also help with the problem of failing to select important variables in stage (1)
- One major challenge with this approach, however, is how exactly we average over results in which a covariate was not included in the model

Averaging over unselected variables

- One conservative remedy is to simply assign $p_j = 1$ whenever $j \notin \mathcal{S}$, the set of selected variables from stage 1
- With this substitution in place, we will have, for each variable, a vector of p -values $p_j^{(1)}, \dots, p_j^{(B)}$, where B is the number of random splits, which we can aggregate in a variety of ways
- For the results that follow, I used the median

Multiple split approach applied to example data



As with the semi-penalized LRT, 5 “A” variables are significant

Remarks

- Certainly, the results are much more stable if we average across sample splits
- The other downside, however, (loss of power from splitting the sample in two) cannot be avoided
- It is possible to extend this idea to obtain confidence intervals as well by inverting the hypothesis tests, although the implementation gets somewhat complicated

TCGA data

- To get a feel for how conservative this approach is, let's apply it to the TCGA data ($n = 536$, $p = 17,322$)
- Using the multiple-splitting approach, only a single variable is significant with $p < 0.05$
- This is similar to the semi-penalized LRT, but again in sharp contrast to the marginal results

Stability selection

- One could argue that trying to obtain a classical p -value isn't really the right goal, that what makes sense for single hypothesis testing isn't relevant to high-dimensional modeling
- Consider, then, the idea of *stability selection* (Meinshausen & Bühlmann, 2010), in which we decide that a variable is significant if it is selected in a high proportion of penalized regression models that have been applied to “perturbed” data
- The most familiar way of perturbing a data set is via resampling (i.e., bootstrapping), although the authors also considered other ideas

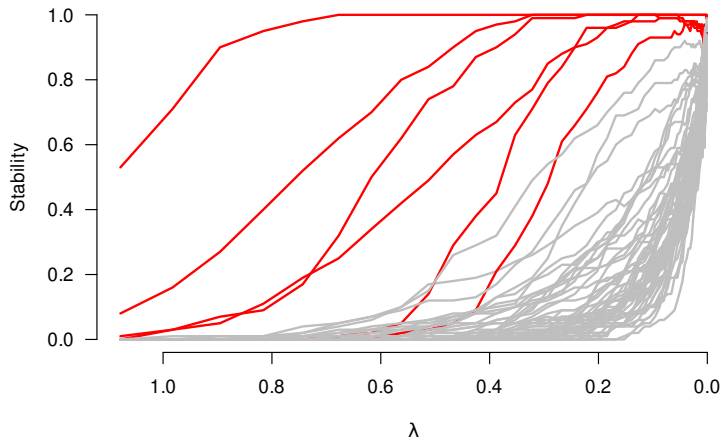
Details

- Furthermore, there are a variety of ways of carrying out bootstrapping, a point we will return to later
- For simplicity, I'll stick to what the authors chose in their original paper: randomly select $n/2$ indices from $\{1, \dots, n\}$ without replacement (this is known as “subagging” and based on an argument that sampling $n/2$ without replacement is fairly similar to resampling n with replacement)
- Letting π_{thr} denote a specified cutoff and $\hat{\pi}_j(\lambda)$ the fraction of times variable j is selected for a given value of λ , the set of *stable variables* is defined as

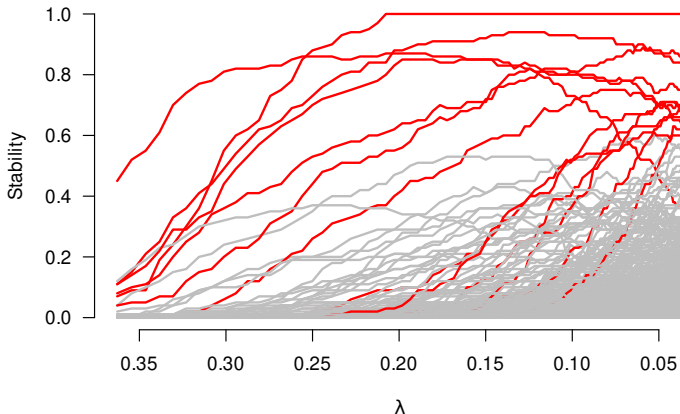
$$\{j : \hat{\pi}_j(\lambda) > \pi_{\text{thr}}\}$$

Stability selection for example data

Variables with $\beta_j \neq 0$ in red:



Stability selection for TCGA data



13 variables exceed $\pi_{\text{thr}} = 0.6$ for any λ (in red)

FDR bound

- Meinshausen & Bühlmann also provide an upper bound for the expected number of false selections in the stable set (i.e., variables with $\beta_j = 0$ and $\hat{\pi}_j(\lambda) > \pi_{\text{thr}}$):

$$\frac{1}{2\pi_{\text{thr}} - 1} \frac{S(\lambda)^2}{p},$$

where $S(\lambda)$ is the expected number of selected variables

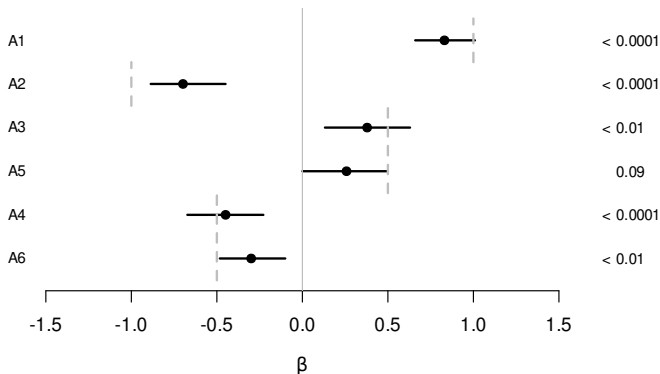
- Note that this bound can only be applied if $\pi_{\text{thr}} > 0.5$
- In practice, however, this bound is rather conservative:
 - For the example data set, only the two variables with $\beta_j = 1$ can be selected at an FDR of 10%
 - For the TCGA data set, only two variables can be stably selected

Bootstrapping

- Stability selection is essentially just bootstrapping, with a special emphasis on whether $\hat{\beta}_j^{(b)} = 0$
- There are a variety of ways of carrying out bootstrapping for regression models; the one we have just seen, in which one selects random elements from $\{(\mathbf{x}_i, y_i)\}_{i=1}^n$, is known as the *pairs bootstrap* or *pairwise bootstrap*
- Alternative methods, such as bootstrapping the residuals, are somewhat less robust in the presence of model uncertainty, as they depend on the selection of an initial model

Bootstrap intervals: Example data

Bootstrap percentile intervals for the six coefficients with $\beta_j \neq 0$, residual approach, λ fixed at $\hat{\lambda}_{CV}$



Does bootstrapping work?

- This is interesting, but a natural question would be whether or not bootstrapping actually works in this setting
- In particular, we have theoretical results establishing that bootstrapping works for maximum likelihood; do those proofs extend to penalized likelihood settings?
- The answer is “no”, or at least, not in the classical sense

Limitations/failures of bootstrapping

There are two fundamental issues with the bootstrap

- One is caused by the sparsity of the lasso solutions; if $\hat{\beta}_j = 0$ in most bootstrap samples, the resulting $(0, 0)$ quantile interval is clearly problematic
- The other is caused by the systematic shrinkage of the lasso estimates; it can be shown that even asymptotically, this never goes away for the lasso (although it does for MCP and SCAD)

Bootstrap intervals for MCP

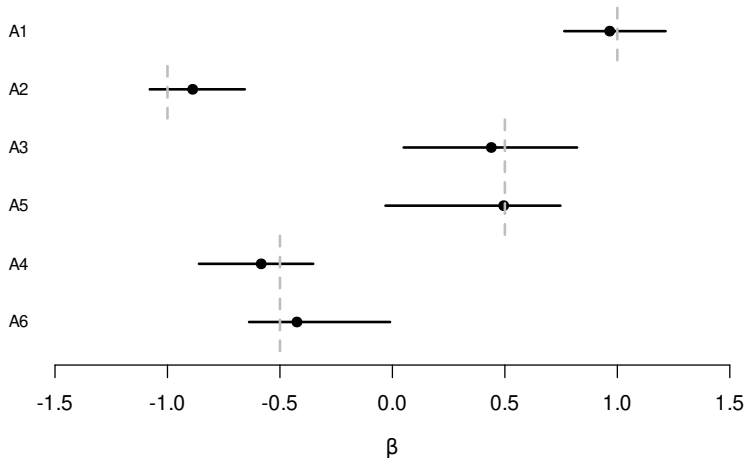


Figure from Harris & Breheny (2025)

Once the sparse draw problem is corrected, however, bootstrap intervals do have correct average coverage:

