

# Making Good Prediction

## A Theoretical Framework

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## Prediction is important

- Predictions of violent events: civil wars, revolutions, local conflicts



## Prediction is important

- Predictions of oncoming recessions



## Prediction is important

- GWAS-level predictions of disease-status



# Motivation

Prediction used in causal inference techniques

- ▶ Selecting amongst many and possibly weak instrumental variables in 2SLS (Belloni et al. 2014)
- ▶ Creating synthetic controls in DID designs (Xu 2015)
- ▶ Collecting interactions in conjoint analysis using LASSOplus (Ratkovic & Tingley 2015)
- ▶ Collecting fewer interactions of factors in factorial experiments using LASSO (Egami & Imai)
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**Not nearly enough attention on creating a framework from which to theoretically consider predictivity**

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  - ▶ There may be logic to and benefits from creation of prediction measures from a prediction framework



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  - ▶ Why is it that certain approaches perform better than others in some scenarios can be hard to ascertain; what is the benchmark against which to compare?
- ▶ Both approaches additionally suffer from curse of dimensionality constraints vis a vis joint/interactive variables as variable size grows.

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- ▶ **Identify sample-appropriate measures for measuring predictivity that match the theoretical solution**
  - ▶ Solution doesn't actually have usable sample analog form. Our second major contribution stems from considering an alternative solution with a sample analog that is useable.

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- ▶ Reflect predictive power of a given variable set
- ▶ Handle groups of variables
- ▶ Be able to differentiate between truly influential variables and noisy variables

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Imagine now we have many  $X$  variables to consider. Then what we are looking for is the set of  $X$  variables that maximize the correct prediction rate,  $c$ .

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$$c = c[p_{\mathbf{X}_d}, p_{\mathbf{X}_c}] = 1 - e[p_{\mathbf{X}_d}, p_{\mathbf{X}_c}] = \frac{1}{2} \sum_{x \in \Pi_x} \max\{p_{\mathbf{X}_d}(x), p_{\mathbf{X}_c}(x)\}$$

$$c[p_{\mathbf{X}_d}, p_{\mathbf{X}_c}] = \frac{1}{2} + \frac{1}{4} \sum_{x \in \Pi_x} |p_{\mathbf{X}_d}(x) - p_{\mathbf{X}_c}(x)| \quad (1)$$

# Problems with Sample Analog

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Sample analog of equation (1) is always increasing in variables and favors ever-increasing the variable set with both truly influential as well as noisy and un-influential variables.

# Suggested Alternative Solution

## Lemma 1

Let  $a_1, a_2, a_3 \dots a_k$  be  $k$  nonnegative numbers. Then

$\sum_{i=1}^k a_i \geq \sqrt{\sum_{i=1}^k a_i^2}$ . If we replace  $a_i$  by  $|p(i|d) - p(i|c)| \forall i, 1 \leq i \leq k$ ,

it is clear that by maximizing  $\sum_{i=1}^k (p_i(d) - p_i(c))^2$  over possible pairs will have the parallel effect of encouraging selection of probability pairs that satisfy the maximization in Equation 1, yielding a better predictor.

We can show that the  $I$ -score can be seen asymptotically as precisely the maximization of the term up to a constant

$$A(\pi_x) = \sum_{i=1}^k (p_i(d) - p_i(c))^2.$$

Since  $\sum_{i=1}^k |p_i(d) - p_i(c)| \geq \sqrt{\sum_{i=1}^k (p_i(d) - p_i(c))^2}$ , a strategy that seeks for a variable set with larger value of  $A(\pi_x)$  will automatically have the effect of seeking for the variable set with a better prediction rate.

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- ▶  $n$  observations of  $Y$  and large number  $S$  of  $\mathbf{X}$ s,  $X_1, X_2, \dots, X_S$ .
- ▶ Randomly select small group,  $m$ , of the  $\mathbf{X}$ s. Call this  $m$   $X_j$ ,  $j = 1, \dots, m$  that take values 0, 1, and 2 (here, discrete example)

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- ▶  $I$ -score designed to place greater weight on cells with more observations:

$$I_{\Pi} = \sum_{k=1}^{m_1} \frac{n_k}{n} \cdot \frac{(\bar{Y}_k - \bar{Y})^2}{\frac{s^2}{n_k}} = \frac{\sum_{k=1}^{m_1} n_k^2 (\bar{Y}_k - \bar{Y})^2}{\sum_{i=1}^n (Y_i - \bar{Y})^2} \quad (2)$$

where  $s^2 = \frac{1}{n} \sum_{i=1}^n (Y_i - \bar{Y})^2$ .

# I-score

- ▶ Asymptotics
- ▶ Data simulations

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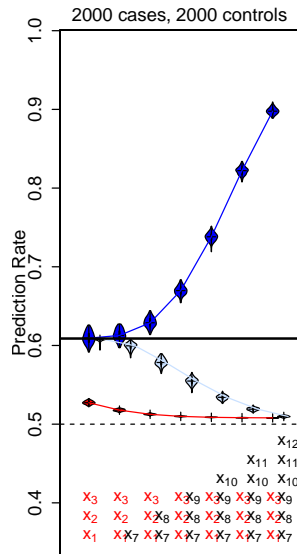
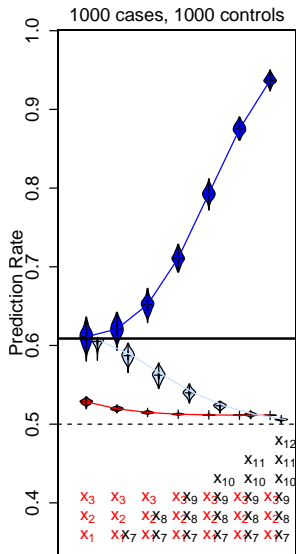
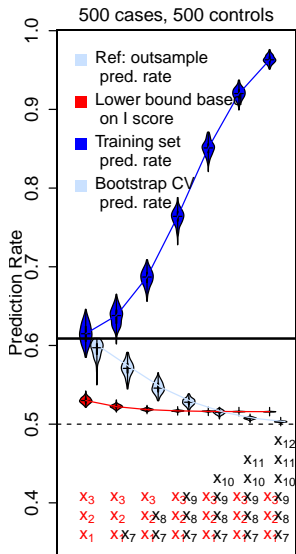
I-score asymptotics

# I-score with data

How does I-score fare with data (sample constrained world)?



# I-score with simulated data



## I-score with real data

	Systematic name	Gene name	Marginal p-value
1	Contig45347_RC	KIAA1683	0.008
2	NM_005145	GNG7	0.54
3	Z34893	ICAP-1A	0.15
4	NM_006121	KRT1	0.9
5	NM_004701	CCNB2	0.003
	<b>Joint I-score: 2.89</b>	<b>Joint p-value: 0.005</b>	Family-wise threshold: $6.98 \times 10^{-5}$

**Table: Real data example** vant Veer

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- ▶ **No model specification:** Requires no specification of a model for joint effect of  $\{X_1, X_2, \dots, X_m\}$  on  $Y$ .  $I$  captures discrepancy between the conditional means of  $Y$  on  $\{X_1, X_2, \dots, X_m\}$  and mean of  $Y$ .

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- ▶ **Differentiation between noisy and influential variables:**  $I$  doesn't monotonically increase with the addition of any variables (as would the sample analog form of Eqn 1). Rather, given a variable set of size  $m$  with  $m - 1$  truly influential variables, the  $I$  is higher under the influential  $m - 1$  variables than under all  $m$  variables. Dropping to  $m - 2$  variables leads to decrease in  $I$ .  $I$  has natural tendency to "peak" at variable set(s) that lead to the maximum predictive rate in the face of noisy variables, under the current sample size.

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# I-score Asymptotics

As  $n \rightarrow \infty$ , it can be shown that the I-score decomposes to two terms that converge to 0 in probability and a third term, call it  $B_n$  that approximates Equation 1 (correct prediction rate) via Lemma 1:

$$\frac{B_n}{n^2} = \lambda^2(1 - \lambda)^2 \sum_{j \in \Pi} [p(j|d) - p(j|c)]^2$$

(Where  $\lim_n \frac{n_d}{n} = \lambda$ , a fixed constant between 0 and 1)

Ignoring the constant term above, the I-score is exactly trying to search for the  $X$  partitions which maximize the summation term

$\sum_{j \in \Pi} [p(j|d) - p(j|c)]^2$ . Recall  $c$ :

$$c[p_{\mathbf{X}_d}, p_{\mathbf{X}_c}] = \frac{1}{2} + \frac{1}{4} \sum_{x \in \Pi_x} |p_{\mathbf{X}_d}(x) - p_{\mathbf{X}_c}(x)| \quad (4)$$

[back](#)