March 23 Recitation

• BIC Score
$$BIC = L(x; \hat{\theta}) - \frac{k}{2} \log(n)$$

$$\begin{cases} k: num. of params. \\ n: num of samples \\ \hat{\theta}: maximum likelyhard estimates of params \end{cases}$$

Regression tree example:

Model I: expr is not factor specific

$$\Rightarrow \mathcal{L}(X;\theta) = \log \frac{m}{m} \frac{\pi}{m} N(X_{it}; M, 6^{2}) = \sum_{t=1}^{m} \sum_{i=1}^{h} \left[-\frac{1}{2} \log_{2}\pi \delta^{2} - \frac{1}{2\delta_{2}} (X_{it} - M)^{2} \right]$$

$$MLE: \text{ find } \hat{\theta} = \operatorname{argmax} \mathcal{L}(X;\theta) = \left\{ (M, 6) \mid \frac{\partial \mathcal{L}(X;\theta)}{\partial M} = \frac{\partial \mathcal{L}(X;\theta)}{\partial \delta} = 0 \right\}$$

$$\text{(maximum likely hord)}$$

$$\Rightarrow \hat{M} = \frac{1}{n \text{ in}} \sum_{t=1}^{m} \sum_{i=1}^{h} X_{it}^{2}$$

$$\hat{\theta} = \left(\frac{1}{n \text{ in}} \sum_{t=1}^{m} \sum_{i=1}^{h} (X_{it} - \hat{M})^{2} \right)^{\frac{1}{2}}$$

$$\Rightarrow \mathcal{L}(X;\hat{\theta}) = n \text{ in } \left(-\frac{1}{2} - \frac{1}{2} \log_{2}(2\pi \hat{\theta}^{2}) \right)$$

$$\Rightarrow BIC = n \text{ in } \left(-\frac{1}{2} - \frac{1}{2} \log_{2}(2\pi \hat{\theta}^{2}) \right) - \frac{2}{2} \log_{2}(n \text{ in })$$

Model 2: Expr. depends on one factor w/ one threshold

Assume it's factor
$$j$$
. For a given threshold f_{j}^{*} $\left\{\begin{array}{c} I_{1}=2t:\ f_{j}t< f_{j}^{*}\end{array}\right\}$

Find Mi, bi, Mi, bi in a similar wary

Note: also need to optimize over all possible thresholds fix

$$\Rightarrow BIC = n|I_1| \left(-\frac{1}{z} - \frac{1}{z} \log(2\pi \hat{\delta}_1^2) \right) + n|I_2| \left(-\frac{1}{z} - \frac{1}{z} \log(2\pi \hat{\delta}_2^2) \right) - \frac{5}{z} \log(nm)$$

- Bayesian model selection
 - > Compare model | and model 2 in general, rather than

best (model 1) vs. best (model 2) defined by MLE

- > Take the distribution of parameter into account
- ⇒ Avoids overfitting (Bayesian Oceam's Razor)

assume PLM) ~ uniform

marginal likely hovel / evidence

Complex model might not always have the best P(DIM)

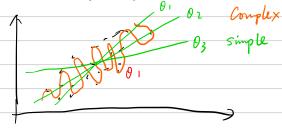
5 plo'/M)=1 - Complex model can get very good P(D/M,0) but only Conservation of prob. mass

(assuming model M)

for very few 0

Simple model might get desent PIDIM, 0) for a large range

 $\theta \mid \theta \Rightarrow \text{Intergral might be biggen}$



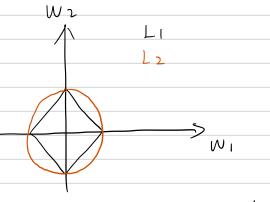
Do: autual data => M2 fits hest Complexity: MI < MZ < M3 (more complex model can model a larger range of D) but resulting in "thiner" distribution

Note: BIC is an approximation of BMS

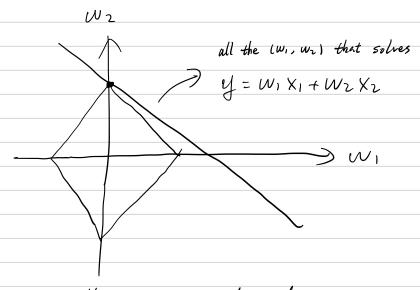
BIC also penalizes complex model (by $\frac{k}{2} \log n$)

•
$$L_1/L_2$$
 penalty
$$L = \sum_{i} (y_i - wx_i)^2 + R$$

$$R = \begin{cases} ||W||_1 = \sum_{k=1}^{d} W_k \\ \frac{1}{2} ||W||_2 = \frac{1}{2} \sum_{k=1}^{d} W_k \end{cases}$$



All the (w, w) with a fixed R



Grandually shrink the diamand until only one point crosses with $y = w_1 x_1 + w_2 x_1 \Rightarrow \text{end up at}$ a wertex

$$0LS \Rightarrow (x^{T}x)^{-1}x^{T}y$$

$$0LS \Rightarrow (x^{T}x + \chi I)^{-1}x^{T}y$$

$$w/L_{2}$$