S&DS 355 / 365 / 565

Data Mining and Machine Learning

## **Model Selection**

Thursday, September 19th

### **Outline for next topics**

SDS 355 won't have to know some of this

- Model selection
- Ridge regression
- The lasso

#### What did we talk about last time?

- Stochastic gradient descent is a first order (first derivatives) method that scales to large classification and regression problems
- Cross validation is a practical way of estimating the variability of test error.

Data: n observations, p predictors

• Use all predictors?

$$y = \beta_0 + \beta_1 x_1 + \dots + \beta_p x_p + \epsilon$$

Data: n observations, p predictors / independent variables

Use all predictors?

$$y = \rho_0 + \rho_1 x_1 + \cdots + \rho_{\rho} x_{\rho} + \epsilon$$

 $y = \beta_0 + \beta_1 x_1 + \dots + \beta_p x_p + \epsilon$  Include Total number of possible subsets of variables to include:  $2^p$ .

Data: n observations, p predictors

Use all predictors?

$$y = \beta_0 + \beta_1 x_1 + \dots + \beta_p x_p + \epsilon$$

- Total number of possible subsets of variables to include: 2<sup>p</sup>.
- Bias-variance tradeoff in number of predictors included.

you don't want to overfit, so we may to reduce the number of parameters in model

#### Data: n observations, p predictors

Use all predictors?

$$y = \beta_0 + \beta_1 x_1 + \dots + \beta_p x_p + \epsilon$$

- Total number of possible subsets of variables to include: 2<sup>p</sup>.
- Bias-variance tradeoff in number of predictors included.
- More complex models are less interpretable.

### Approaches to feature selection

- similar to HW
- Subset selection use a "good subset" of the p predictors
- Shrinkage use all *p* predictors but encourage more coefficients to be near 0
- Dimension reduction condense the set of predictors by projecting to a lower subspace



Options range from null model  $\mathcal{M}_0$  (no predictors) to full model  $\mathcal{M}_p$  containing all p predictors.

• Fit  $\mathcal{M}_0$ .

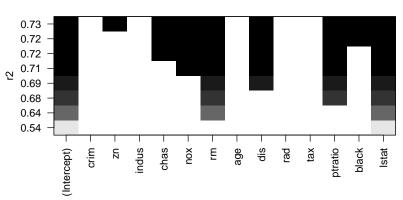
- Fit  $\mathcal{M}_0$ .
- For k = 1, 2, ..., p, identify the best model  $\mathcal{M}_k$  using k of the p predictors judged via training error.

- Fit  $\mathcal{M}_0$ .
- For k = 1, 2, ..., p, identify the best model  $\mathcal{M}_k$  using k of the p predictors judged via training error.
  - e.g. for regression: use RSS, R<sup>2</sup>

- Fit  $\mathcal{M}_0$ .
- For k = 1, 2, ..., p, identify the best model  $\mathcal{M}_k$  using k of the p predictors judged via training error.
  - ▶ e.g. for regression: use RSS, R<sup>2</sup>
  - e.g. for classification: use misclassification error, deviance

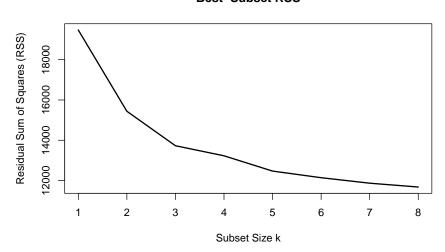
- Fit  $\mathcal{M}_0$ .
- For k = 1, 2, ..., p, identify the best model  $\mathcal{M}_k$  using k of the p predictors judged via training error.
  - ▶ e.g. for regression: use RSS, R<sup>2</sup>
  - e.g. for classification: use misclassification error, deviance
- Select the best model among  $\mathcal{M}_0, \mathcal{M}_1, \dots, \mathcal{M}_p$  on basis of cross-validated prediction error.

```
library(MASS)
library(leaps)
m1 <- regsubsets(medv ~ ., data=Boston, nbest=1, method="exhausting")</pre>
```



#### Boston dataset:





Not feasible when p is large. There are  $2^p$  models to consider!

e.g. 
$$p = 5 \Rightarrow 2^5 = 32$$
 models

$$p = 10 \Rightarrow 2^{10} = 1024 \text{ models}$$

$$p = 100$$
?

Instead of computing all combinations,

1. Forward stepwise selection

Starting from the null model, build an increasing sequence of *nested models*.

greedy selection takes a locally optimal decision at every step but often this works out.

pick the cavariate that gives the best model, and keep daing this in a loop.

1. Forward stepwise selection

Starting from the null model, build an increasing sequence of *nested* models.

- Start with M<sub>0</sub>.
- For k = 1, ..., p, pick the best **one** of the remaining unused predictors to add to  $\mathcal{M}_{k-1}$  to form  $\mathcal{M}_k$ .
- Select the best model among  $\mathcal{M}_0, \mathcal{M}_1, \dots, \mathcal{M}_p$  on basis of estimated prediction error.

2. Backward stepwise selection

Starting from the full model, build a decreasing sequence of *nested models*.

#### 2. Backward stepwise selection

Starting from the full model, build a decreasing sequence of *nested models*.

- Start with M<sub>p</sub>.
- For k = p 1, p 2, ..., 0, pick the worst **one** of the existing predictors to remove from  $\mathcal{M}_{k+1}$  to form  $\mathcal{M}_k$ .
- Select the best model among  $\mathcal{M}_0, \mathcal{M}_1, \dots, \mathcal{M}_p$  on basis of estimated prediction error.
  - nsdel uthu ==0

2. Backward stepwise selection

Starting from the full model, build a decreasing sequence of *nested models*.

- Start with  $\mathcal{M}_p$ .
- For k = p 1, p 2, ..., 0, pick the worst **one** of the existing predictors to remove from  $\mathcal{M}_{k+1}$  to form  $\mathcal{M}_k$ .
- Select the best model among  $\mathcal{M}_0, \mathcal{M}_1, \dots, \mathcal{M}_p$  on basis of estimated prediction error.

Backward and forward stepwise selection are more computationally feasible than best subsets, but no guarantee they'll find the best subset of the p predictors to use.

| because greedy choices are not predicted applicably optimal 1/24

- 3. Bidirectional stepwise selection
  - Start with either the null model or the full model

- 3. Bidirectional stepwise selection
  - Start with either the null model or the full model
  - At each step, consider the impact of adding or subtracting a predictor

- 3. Bidirectional stepwise selection
  - Start with either the null model or the full model
  - At each step, consider the impact of adding or subtracting a predictor
  - Stop when no further improvements can be made

- 3. Bidirectional stepwise selection
  - Start with either the null model or the full model
  - At each step, consider the impact of adding or subtracting a predictor
  - Stop when no further improvements can be made
    - Will not cycle because model must reduce RSS

- 3. Bidirectional stepwise selection
  - Start with either the null model or the full model
  - At each step, consider the impact of adding or subtracting a predictor
  - Stop when no further improvements can be made
    - Will not cycle because model must reduce RSS
    - Will eventually stop because only finitely many models

- 3. Bidirectional stepwise selection
  - Start with either the null model or the full model
  - At each step, consider the impact of adding or subtracting a predictor
  - Stop when no further improvements can be made
    - Will not cycle because model must reduce RSS
    - Will eventually stop because only finitely many models
    - ▶ Could run for more than O(p) steps

• RSS is a bad metric to use (as is multiple R2). (Why?)

- RSS is a bad metric to use (as is multiple R<sup>2</sup>). (Why?)
- Cross-validated MSE is a good criterion, but is time consuming.

- RSS is a bad metric to use (as is multiple R<sup>2</sup>). (Why?)
- Cross-validated MSE is a good criterion, but is time consuming.
- Other options?

- Mallow's  $C_p$  (regression only)
- AIC (regression or classification)
- BIC (regression or classification)

$$C_p = \frac{1}{n}(RSS + 2p\widehat{\sigma}^2),$$

where p is the number of coefficients fitted and  $\hat{\sigma}^2$  is estimated error variance.

$$C_p = \frac{1}{n}(RSS + 2p\widehat{\sigma}^2),$$

where p is the number of coefficients fitted and  $\hat{\sigma}^2$  is estimated error variance. Derivation Setup:

Data: (X, Y), X is  $n \times p$  and Y is  $n \times 1$ 

Derivation not needed

Fitted model:  $\widehat{Y} = X\widehat{\beta}$ 

Consider how well our model predicts (X, Y), measured via out-of-sample MSE:

$$MSE_{OOS} = E\left[\frac{1}{n}\sum_{i=1}^{n}(\widetilde{Y}_{i}-\widehat{Y}_{i})^{2}\right]$$

$$MSE_{OOS} = E\left[\frac{1}{n}\sum_{i=1}^{n}(\widetilde{Y}_{i}-\widehat{Y}_{i})^{2}\right]$$

For any i,

$$E[(\widetilde{Y}_i - \widehat{Y}_i)^2] = Var(\widetilde{Y}_i - \widehat{Y}_i) + (E[\widetilde{Y}_i - \widehat{Y}_i])^2$$

$$= Var(\widetilde{Y}_i) + Var(\widehat{Y}_i) - 2Cov(\widetilde{Y}_i, \widehat{Y}_i) + [E(\widetilde{Y}_i) - E(\widehat{Y}_i)]^2$$

$$MSE_{OOS} = E\left[\frac{1}{n}\sum_{i=1}^{n}(\widetilde{Y}_{i}-\widehat{Y}_{i})^{2}\right]$$

For any i,

$$E[(\widetilde{Y}_i - \widehat{Y}_i)^2] = Var(\widetilde{Y}_i - \widehat{Y}_i) + (E[\widetilde{Y}_i - \widehat{Y}_i])^2$$

$$= Var(\widetilde{Y}_i) + Var(\widehat{Y}_i) - 2Cov(\widetilde{Y}_i, \widehat{Y}_i) + [E(\widetilde{Y}_i) - E(\widehat{Y}_i)]^2$$

Note that  $Cov(\widetilde{Y}_i, \widehat{Y}_i) = 0$ , so:

$$E[(\widetilde{Y}_i - \widehat{Y}_i)^2] = Var(\widetilde{Y}_i) + Var(\widehat{Y}_i) + [E(\widetilde{Y}_i) - E(\widehat{Y}_i)]^2$$

In-sample (IS) MSE:

$$MSE_{IS} = E\left[\frac{1}{n}\sum_{i=1}^{n}(Y_i - \widehat{Y}_i)^2\right]$$

### Mallow's $C_{D}$

In-sample (IS) MSE:

$$MSE_{IS} = E\left[\frac{1}{n}\sum_{i=1}^{n}(Y_i - \widehat{Y}_i)^2\right]$$

For any i,

$$E[(Y_i - \widehat{Y}_i)^2] = Var(Y_i - \widehat{Y}_i) + (E[Y_i - \widehat{Y}_i])^2$$

$$= Var(Y_i) + Var(\widehat{Y}_i) - 2Cov(Y_i, \widehat{Y}_i) + [E(Y_i) - E(\widehat{Y}_i)]^2$$

In-sample (IS) MSE:

$$MSE_{IS} = E\left[\frac{1}{n}\sum_{i=1}^{n}(Y_i - \widehat{Y}_i)^2\right]$$

For any i,

$$E[(Y_i - \widehat{Y}_i)^2] = Var(Y_i - \widehat{Y}_i) + (E[Y_i - \widehat{Y}_i])^2$$

$$= Var(Y_i) + Var(\widehat{Y}_i) - 2Cov(Y_i, \widehat{Y}_i) + [E(Y_i) - E(\widehat{Y}_i)]^2$$

Note  $Y_i$  and  $\widetilde{Y}_i$ :

- are independent
- have the same distribution, e.g.,  $Var(Y_i) = Var(\widetilde{Y}_i)$  and  $E(Y_i) = E(\widetilde{Y}_i)$

*i*-th term in the summation of MSE<sub>OOS</sub> again:

$$E[(\widetilde{Y}_i - \widehat{Y}_i)^2] = Var(\widetilde{Y}_i) + Var(\widehat{Y}_i) + [E(\widetilde{Y}_i) - E(\widehat{Y}_i)]^2$$

*i*-th term in the summation of  $MSE_{OOS}$  again:

$$E[(\widetilde{Y}_i - \widehat{Y}_i)^2] = Var(Y_i) + Var(\widehat{Y}_i) + [E(Y_i) - E(\widehat{Y}_i)]^2$$
  
= 
$$E[(Y_i - \widehat{Y}_i)^2] + 2Cov(Y_i, \widehat{Y}_i)$$

Averaging over all i, we get:

$$\frac{1}{n}E\left[\sum(\widetilde{Y}_i-\widehat{Y}_i)^2\right]=\frac{1}{n}E\left[\sum(Y_i-\widehat{Y}_i)^2\right]+\frac{2}{n}\sum Cov(Y_i,\widehat{Y}_i)$$

*i*-th term in the summation of *MSE<sub>OOS</sub>* again:

$$E[(\widetilde{Y}_i - \widehat{Y}_i)^2] = Var(Y_i) + Var(\widehat{Y}_i) + [E(Y_i) - E(\widehat{Y}_i)]^2$$
$$= E[(Y_i - \widehat{Y}_i)^2] + 2Cov(Y_i, \widehat{Y}_i)$$

Averaging over all i, we get:

$$\frac{1}{n}E\left[\sum(\widetilde{Y}_i-\widehat{Y}_i)^2\right]=\frac{1}{n}E\left[\sum(Y_i-\widehat{Y}_i)^2\right]+\frac{2}{n}\sum Cov(Y_i,\widehat{Y}_i)$$

We can show that  $\sum Cov(Y_i, \widehat{Y}_i) = \sigma^2 p$ .

*i*-th term in the summation of *MSE<sub>OOS</sub>* again:

$$E[(\widetilde{Y}_i - \widehat{Y}_i)^2] = Var(Y_i) + Var(\widehat{Y}_i) + [E(Y_i) - E(\widehat{Y}_i)]^2$$
$$= E[(Y_i - \widehat{Y}_i)^2] + 2Cov(Y_i, \widehat{Y}_i)$$

Averaging over all i, we get:

$$\frac{1}{n}E\left[\sum(\widetilde{Y}_i-\widehat{Y}_i)^2\right]=\frac{1}{n}E\left[\sum(Y_i-\widehat{Y}_i)^2\right]+\frac{2}{n}\sum Cov(Y_i,\widehat{Y}_i)$$

We can show that  $\sum Cov(Y_i, \widehat{Y}_i) = \sigma^2 p$ .

In summary,

$$MSE_{OOS} = MSE_{IS} + \frac{2p\sigma^2}{n}$$

We approximate  $MSE_{IS}$  using RSS/n and  $\sigma^2$  using  $\widehat{\sigma}^2$ .

$$C_p = \frac{RSS}{n} + \frac{2p\widehat{\sigma}^2}{n}.$$

- Adjusts RSS with a penalty that depends on number predictors and variance of error term.
- If  $\hat{\sigma}^2$  is unbiased estimate of  $\sigma^2$ , then  $C_p$  is an unbiased estimate of test MSE.

We approximate  $MSE_{IS}$  using RSS/n and  $\sigma^2$  using  $\widehat{\sigma}^2$ .

$$C_p = \frac{RSS}{n} + \frac{2p\widehat{\sigma}^2}{n}.$$

- Adjusts RSS with a penalty that depends on number predictors and variance of error term.
- If  $\hat{\sigma}^2$  is unbiased estimate of  $\sigma^2$ , then  $C_p$  is an unbiased estimate of test MSE.

To summarize, choose model with lowest  $C_p$ .

#### **AIC**

Akaike Information Criterion (regression or classification):

$$AIC = -2\log L + 2p,$$

where L is the likelihood of the model.

We can show for linear regression,

$$-2\log L = \frac{RSS}{\widehat{\sigma}^2} + C,$$

for some constant C. Hence,

$$AIC = \frac{1}{n\widehat{\sigma}^2}(RSS + 2p\widehat{\sigma}^2).$$

Very similar to  $C_p$ 

#### **BIC**

Bayesian Information Criterion (regression or classification):

$$BIC = -2\log L + p\log(n).$$

For regression,

$$BIC = \frac{1}{n}(RSS + \log(n)p\widehat{\sigma}^2).$$

How do AIC and BIC compare?

- Penalty on AIC: 2p
- Penalty on BIC: log(n)p

#### **BIC**

Bayesian Information Criterion (regression or classification):

$$BIC = -2\log L + p\log(n).$$

For regression,

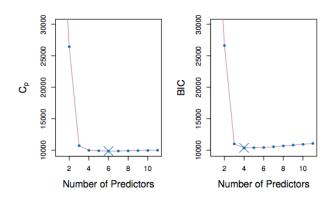
$$BIC = \frac{1}{n}(RSS + \log(n)p\widehat{\sigma}^2).$$

How do AIC and BIC compare?

- Penalty on AIC: 2p
- Penalty on BIC: log(n)p
- $\log(n) > 2$  for n > 7

BIC has heavier penalty on number of variables, produces smaller models.

## Comparison



#### **Summary**

- We like models that minimize expected test error.
- Cross-validation is nice, but requires a lot of computation.
- Stepwise model selection allows us to pick a model based on some measure of expected test error using in-sample measures like  $C_p$ , AIC, or BIC.

### Next up: Shrinkage and selection

An alternative to variable selection is to simply use all predictors, but impose a penalty on the magnitude of the coefficients.

Ridge regression

Result is that the coefficients get *shrunken* towards 0 and standard error of coefficients is much lower.

The Lasso combines shrinkage and selection.