# CPSC 340: Machine Learning and Data Mining

Feature Selection Fall 2017

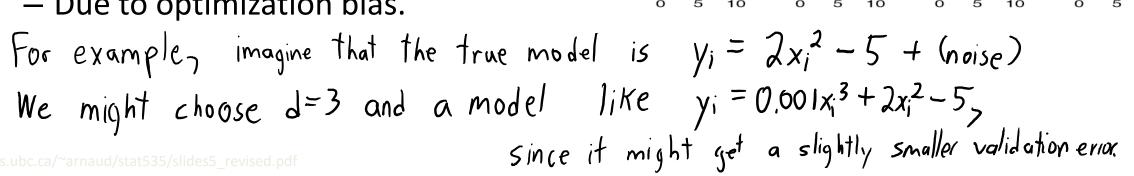
#### Admin

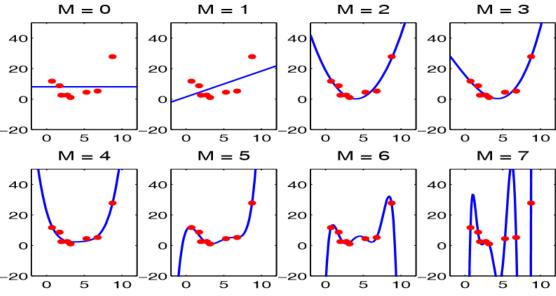
- Assignment 2:
  - 1 late day to hand in tonight, 2 for Wednesday, answers posted Thursday.
- Extra office hours
  - Thursday at 4pm (ICICS 246).
- Midterm details:
  - Friday in class, details on Piazza.

# Last Time: Finding the "True" Model

- What if y<sub>i</sub> really is a polynomial function of x<sub>i</sub>?
  - How can we find the "true" degree 'p' of the polynomial?

- Training error does not work:
  - It goes down as 'p' goes up.
- Cross-validation may also not work:
  - Tends to overestimate 'p'.
  - Due to optimization bias.





### Last Time: Complexity Penalties

- We discussed putting a penalty on the model complexity.
  - Want to fit the data and have a simple model.
- For example, minimize training error plus the degree of polynomial.

- If we use p=4, use "training error plus 4" as error.
- If two 'p' values have similar error, this prefers the smaller 'p'.
  - Can't optimize this using gradient descent, since it's discontinuous in 'p'.

# Choosing Degree of Polynomial Basis

How can we optimize this score in terms of 'w' and 'p'?

$$Score(w_{2}p) = \frac{1}{2}||Z_{pw} - y||^{2} + p$$

- Form  $Z_2$ , solve for 'w', compute  $\frac{1}{2}||Z_2w-y||^2$ .
  - Stop if it doesn't improve by at least 1.
- Form  $Z_3$ , solve for 'w', compute  $\frac{1}{2}||Z_3w-y||^2$ .
  - Stop if it doesn't improve by at least 1.
- The "at least 1" criteria makes sure increasing 'p' has a non-trivial improvement.

- Form 
$$Z_0$$
, solve for 'w', compute  $\frac{1}{2}||Z_0w-y||^2$ .  
- Form  $Z_1$ , solve for 'w', compute  $\frac{1}{2}||Z_1w-y||^2$ .  
• Stop if it doesn't improve by at least 1.  
- Form  $Z_2$ , solve for 'w', compute  $\frac{1}{2}||Z_2w-y||^2$ .  
• Stop if it doesn't improve by at least 1.  
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### Information Criteria

There are many scores, usually with the form:

$$Score(w_{2}p) = \frac{1}{2}||Z_{pw} - y||^{2} + \lambda p$$

- The new parameter  $\lambda > 0$  controls how strong we penalize complexity.
  - "You need to decrease the training error by least  $\lambda$  to increase 'p' by 1".
- Previous case ( $\lambda = 1$ ) is called Akaike information criterion (AIC).
- Other choices of  $\lambda$  give other criteria:
  - Mallow's C<sub>p</sub>.
  - Adjusted R<sup>2</sup>.

# **Bayesian Information Criterion**

- A disadvantage of these methods:
  - Still prefers a larger 'p' as 'n' grows.
- Solution: make λ depend on 'n'.
- For example, the Bayesian information criterion (BIC) uses:

$$\lambda = \frac{1}{2} \log(n)$$

- BIC penalizes a bit more than AIC for large 'n'.
  - As 'n' goes to ∞, recovers "true" model ("consistent" for model selection).
- In practice, we usually just try a bunch of different  $\lambda$  values.
  - Picking  $\lambda$  is like picking 'k' in k-means.

#### Discussion of other Scores for Model Selection

- There are many other scores:
  - Elbow method (similar to choosing  $\lambda$ ).
    - You could also use BIC for choosing 'k' in k-means.
  - Methods based on validation error.
    - "Take smallest 'p' within one standard error of minimum cross-validation error".
  - Minimum description length.
  - Risk inflation criterion.
  - False discovery rate.
  - Marginal likelihood (CPSC 540).
- These can adapted to use the L1-norm and other errors.

(pause)

### Motivation: Discovering Food Allergies

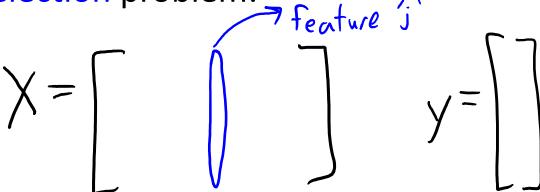
Recall the food allergy example:

Egg	Milk	Fish	Wheat	Shellfish	Peanuts	•••	Sick?
0	0.7	0	0.3	0	0		1
0.3	0.7	0	0.6	0	0.01		1
0	0	0	0.8	0	0		0
0.3	0.7	1.2	0	0.10	0.01		1

- Instead of predicting "sick", we want to do feature selection:
  - Which foods are "relevant" for predicting "sick".

#### Feature Selection

• General feature selection problem:



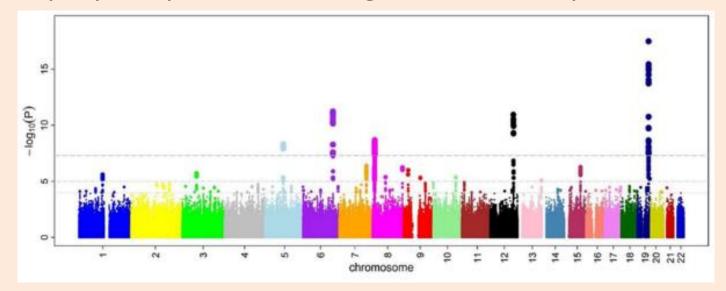
- Find the features (columns) of 'X' that are important for predicting 'y'.
  - "What are the relevant factors?"
  - "What which basis functions should I use among these choices?"
  - "What types of new data should I collect?"
  - "How can I speed up computation?"
- One of most important problems in ML/statistics, but very messy.
  - For now, we'll say a feature is "relevant" if it helps predict  $y_i$  from  $x_i$ .

### "Association" Approach

- A simple/common way to do feature selection:
  - For each feature 'j', compute correlation between feature values and 'y'.
    - Say that 'j' is relevant if correlation is above 0.9 or below -0.9.
- Turns feature selection into hypothesis testing for each feature.
  - There are many other measures of "dependence" (Wikipedia).
- Usually gives unsatisfactory results as it ignores variable interactions:
  - Includes irrelevant variables: "Taco Tuesdays".
    - If tacos make you sick, and you often eat tacos on Tuesdays, it will say "Tuesday" is relevant.
  - Excludes relevant variables: "Diet Coke + Mentos Eruption".
    - Diet coke and Mentos don't make you sick on their own, but together they make you sick.

#### Genome-Wide Association Studies

- Genome-wide association studies:
  - Measure if there exists a dependency between each individual "single-nucleotide polymorphism" in the genome and a particular disease.



- Has identified thousands of genes "associated" with diseases.
  - But by design this has a huge numbers of false positives (and many false negatives).

# "Regression Weight" Approach

- A simple/common approach to feature selection:
  - Fit regression weights 'w' (maybe with least squares).
  - Take all features 'j' where weight |wj| is greater than a threshold.

- This should recognize that "Tuesday" is irrelevant.
  - If you get enough data, and you sometimes eat tacos on other days.
     (And the relationship is actually linear.)
- This should recognize that "Diet Coke" and "Mentos" are relevant.
  - Assuming this combination occurs enough times in the data.

# "Regression Weight" Approach

- A simple/common approach to feature selection:
  - Fit regression weights 'w' (maybe with least squares).
  - Take all features 'j' where weight |wj| is greater than a threshold.

- Has major problems with collinearity:
  - If the "Tuesday" variable always equals the "taco" variable,
     it could say that Tuesdays are relevant but tacos are not.

If you have two copies of an irrelevant feature,
 it could take both irrelevant copies.

#### Search and Score Methods

- Most common feature selection framework is search and score:
  - 1. Define score function f(S) that measures quality of a set of features 'S'.
  - 2. Now search for the variables 'S' with the best score.
- Example with 3 features:
  - Compute "score" of using feature 1.
  - Compute "score" of using feature 2.
  - Compute "score" of using feature 3.
  - Compute "score" of using features {1,2}.
  - Compute "score" of using features {1,3}.
  - Compute "score" of using features {2,3}.
  - Compute "score" of using features {1,2,3}.
  - Compute "score" of using features {}.
  - Return the set of features 'S' with the best "score".

#### Which Score Function?

- The score can't be the training error.
  - Training error goes down as you add features, so will select all features.
- A more logical score is the validation error.
  - "Find the set of features that gives the lowest validation error."
  - To minimize test error, this is what we want.
- But there are problems due to the large number of sets of variables:
  - If we have 'd' variables, there are 2<sup>d</sup> sets of variables.
  - Optimization bias is high: we're optimizing over 2<sup>d</sup> models (not 10).
  - Prone to false positives: irrelevant variables will sometimes help by chance.

### "Number of Features" Penalties

To reduce false positives, we can again use complexity penalties:

$$Score(S) = \frac{1}{2} \sum_{i=1}^{n} (w_s^T x_{is} - y_i)^2 + Size(S)$$

- E.g., we could use squared error and number of non-zeroes.
- We're using ' $x_{is}$ ' as the features 'S' of example  $x_i$ .
- If two 'S' have similar error, this prefers the smaller set.
  - It prefers having  $w_3 = 0$  instead of  $w_3 = 0.00001$ .

• Instead of "size(S)", we usually write this using the "LO-norm"...

### LO-Norm and "Number of Features We Use"

• In linear models, setting  $w_i = 0$  is the same as removing feature 'j':

$$y_{i} = w_{i} x_{i1} + w_{2} x_{i2} + w_{3} x_{i3} + \cdots + w_{d} x_{id}$$

$$y_{i} = w_{i} x_{i1} + 0 + w_{3} x_{i3} + \cdots + w_{d} x_{id}$$

$$y_{i} = w_{i} x_{i1} + 0 + w_{3} x_{i3} + \cdots + w_{d} x_{id}$$
ignore  $x_{i2}$ 

• The LO "norm" is the number of non-zero values.

If 
$$W = \begin{bmatrix} 1 \\ 0 \\ 2 \\ 0 \\ 3 \end{bmatrix}$$
 then  $||w||_0 = 3$  If  $w = \begin{bmatrix} 6 \\ 0 \\ 0 \\ 0 \end{bmatrix}$  then  $||w||_0 = 0$ .

- Not actually a true norm.
- If 'w' has a small L0-norm, then it doesn't use many features.

### L0-penalty: optimization

L0-norm penalty for feature selection:

$$s(ore(w) = \frac{1}{2} || \chi_w - y ||^2 + \frac{1}{2} || w||_0$$

- Suppose we want to use this to evaluate the features S = {1,2}:
  - First fit the 'w' just using features 1 and 2.
  - Now compute the training error with this 'w' and features 1 and 2.
  - Add  $\lambda$ \*2 to the training error to get the score.
- We repeat this with other choices of 'S' to find the "best" features.

### L0-penalty: interpretation

L0-norm penalty for feature selection:

$$s(ore(w) = \frac{1}{2} || \chi_w - y ||^2 + \frac{1}{2} || w||_0$$

- Balances between training error and number of features we use.
  - With  $\lambda$ =0, we get least squares with all features.
  - With  $\lambda$ =∞, we must set w=0 and not use any features.
  - With other  $\lambda$ , balances between training error and number of non-zeroes.
    - Larger  $\lambda$  puts more emphasis on having zeroes in 'w' (more feature selection).
    - Different values give AIC, BIC, and so on.

# Forward Selection (Greedy Search Heuristic)

- In search and score, it's also just hard to search for the best 'S'.
  - There are 2<sup>d</sup> possible sets.

- A common greedy search procedure is forward selection:
  - 1. Start with an empty set of features, S = [].
  - 2. For each possible feature 'j':
    - Compute scores of features 'S' combined with features 'j'.
  - 3. If no 'j' improves the score, stop.
  - 4. Otherwise, add the 'j' that improves the score the most to 'S'.
    - Then go back to Step 2.

# Forward Selection (Greedy Search Heuristic)

- In search and score, it's also just hard to search for the best 'S'.
  - There are 2<sup>d</sup> possible sets.
- A common greedy search procedure is forward selection:

```
1. Compute score if we use no features.

2. Try adding "taco", "milt", "egg", and so on (computing score of each)

3. Add "milk" because it got the best score.

4. Try {milk, taco}, {milk, egg}, and so on (computing score of each

5. Continue until no single-variable addition improves the score.
```

- Not guaranteed to find the best set, but reduces many problems:
  - Considers O(d²) models: cheaper, ovefits less, has fewer false positives.

### Summary

- Information criteria are scores that penalize number of parameters.
  - When we want to find the "true" model.
- Feature selection is task of choosing the relevant features.
  - Obvious simple approaches have obvious simple problems.
- Search and score: find features that optimize some score.
  - L0-norm penalties are the most common scores.
  - Forward selection is a heuristic to search over a smaller set of features.

Next time: getting a good test error even with irrelevant features.

#### Backward Selection and RFE

- Forward selection often works better than naïve methods.
- A related method is backward selection:
  - Start with all features, remove the one that most improves the score.
- If you consider adding or removing features, it's called stagewise.
- Stochastic local search is a class of fancier methods.
- Recursive feature elimination is another related method:
  - Fit parameters of a regression model.
  - Prune features with small regression weights.
  - Repeat.

# Complexity Penalties for Other Models

- Scores like AIC and BIC can also be used in other contexts:
  - When fitting a decision tree, only split a node if it improves BIC.
  - This makes sense if we're looking for the "true tree", or maybe just a simple/interpretable tree that performs well.
- In these cases we replace "L0-norm" with "degrees of freedom".
  - In linear models fit with least squares, degrees of freedom is number of non-zeroes.
  - Unfortunately, it is not always easy to measure "degrees of freedom".