10601 Machine Learning

Model and feature selection

Occam's Razor



- William of Ockham (1285-1349) Principle of Parsimony:
 - "One should not increase, beyond what is necessary, the number of entities required to explain anything."
- Regularization penalizes for "complex explanations"
- Alternatively (but pretty much the same), use *Minimum Description Length (MDL) Principle*:
 - minimize length(misclassifications) + length(hypothesis)
- *length*(misclassifications) e.g., #wrong training examples
- *length*(hypothesis) e.g., size of decision tree

Minimum Description Length Principle

• MDL prefers small hypothesis that fit data well:

$$h_{MDL} = \arg\min_{h} L_{C_1}(\mathcal{D} \mid h) + L_{C_2}(h)$$

- L_{C1}(D|h) description length of data under code C₁ given h
 - Only need to describe points that *h* doesn't explain (classify correctly)
- L_{C2}(h) description length of hypothesis h
- Decision tree example
 - L_{C1}(D|h) #bits required to describe data given h
 - If all points correctly classified, $L_{C1}(D|h) = 0$
 - L_{C2}(h) #bits necessary to encode tree
 - Trade off quality of classification with tree size

What you need to know about Model Selection, Regularization and Cross Validation

Cross validation

- (Mostly) Unbiased estimate of true error
- LOOCV is great, but hard to compute
- *k*-fold much more practical
- Use for selecting parameter values!
- Model selection
 - Search for a model with low cross validation error
- Regularization
 - Penalizes for complex models
 - Select parameter with cross validation
 - Really a Bayesian approach
- Minimum description length
 - Information theoretic interpretation of regularization

Bayesian approach

- Start with a simple model
- As data comes, increase the complexity as necessary
- My research area: Nonparametric Bayes
- The complexity of the model is unbounded
- Select the correct complexity from data (posterior)
- For ex: the number of clusters

Feature selection

- Choose an optimal subset from the set of all N features
 - Only use a subset of a possible words in a dictionary
 - Only use a subset of genes
- Why?
- Can we do model selection to solve this? -2^n models

Two approaches: 1. Filter

- Independent of classifier used
- Rank features using some criteria based on their relevance to the classification task
- For example, mutual information:

$$I(X;Y) = \sum_{y \in Y} \sum_{x \in X} p(x,y) \log \left(\frac{p(x,y)}{p_1(x) p_2(y)} \right),$$

 Choose a subset based on the sorted scores for the criteria used

2. Wrapper

- Classifier specific
- Greedy (large search space)
- Initialize F = null set
 - At each step, using cross validation or an information theoretic criteria, choose a feature to add to the subset [training should be done with only features in F + new feature]
 - Add the chosen feature to the subset
- Repeat until no improvement to CV accuracy

Problem Set 4

Q1.3:

• Take derivatives w.r.t to α first then w,b.

Q. 1.6

- Minimize the violations as much as possible.
- Assume C is large but not ∞ .

Q. 2

- Either explain why some algorithm does not work well.
- Or draw the final result of the algorithms.

Q. 3

The contour of the distribution in 2-D:

- Spherical Gaussian: concentric circles
- Diagonal Gaussian: concentric eclipses with axes parallel to the coordinate axes.
- Unrestricted covariance Gaussian: concentric eclipses

Q. 4

Cutting the tree by thresholding

