Lecture 11: Decision Trees

TTIC 31020: Introduction to Machine Learning

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TTI-Chicago

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 - Hinge loss: exported from SVMs; motivated as finding max-margin separator

Review: binary vs. multi-class classification

- We talked about binary classification in class
- Extensions to multi-class classification exist (e.g., softmax classifier on pset2)
- Sometimes has form of training multiple binary "one vs. all" classifiers

Review: sparsity in classification

- Solving dual of SVM problem leads to sparsity in dual parameters (a relatively small number of "support vectors")
- Solving primal problem with subgradient descent can also lead to sparsity (depending on regularization)
 - If classification is correct (perceptron) or correct with large-enough margin (hinge), there is no loss suffered and therefore no parameter update
 - Intuitively: if a feature is zero for all examples for which a parameter update is performed, its weight won't change (amount of sparsity depending on regularization and initialization)

Review: kernels

- We talked about kernels in the context of SVMs
- But kernels can be used any time we can write a machine learning method in terms of dot products between inputs
- SVMs are a good fit for kernels because of sparse solutions (only support vectors are needed, which reduces computational requirements of using kernels)

General "pipeline" of developing a learning algorithm:

• Write down the complete parameterized objective:

$$\min_{\theta} \sum_{i} f(\theta; \mathbf{x}_i, y_i)$$

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$$g_i(\theta, \mathbf{x}_i, y_i) = \begin{cases} g_i^1(\theta, \mathbf{x}_i, y_i) & \text{if } \theta \leq \theta', \\ g_i^2(\theta, \mathbf{x}_i, y_i) & \text{if } \theta > \theta'. \end{cases}$$

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• Implement gradient descent, e.g., stochastic:

$$\theta^{(t+1)} = \theta^{(t)} - \eta_t g_{i(t)}(\theta, \mathbf{x}_{i(t)}, y_{i(t)})$$

Space partition by classifiers

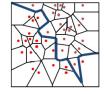
- \bullet We learn a classifier to partition the space of ${\mathcal X}$ according to predicted ${\mathcal Y}$
- Linear classifier: linear partition (hyperplane)



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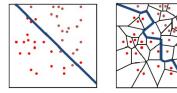


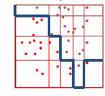


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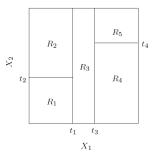
- *k*NN: Voronoi partition data driven, non-linear, no simple parametric form
- Today: decision trees non-linear partition, but with a simple parametric form

Space partition

- Decision boundary: partition the space into regions with fixed prediction value
- Linear classifier: regions = half spaces

Space partition

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- Linear classifier: regions = half spaces
- More general partition: hyper-rectangles

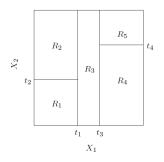


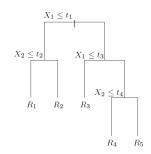
• The regions are easy to describe, e.g.,

$$R_2 = \{ \mathbf{x} : x_1 < t_1 \text{ and } x_2 > t_2 \}$$

Partition tree

 We can describe a non-overlapping partition of the space into hyper-rectangles via a tree:

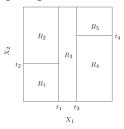


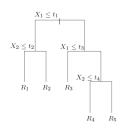


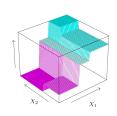
- Regions correspond to leaves
- A point is placed in a region by "dropping" it down the tree, applying a test at each node

Classification and regression trees

- Associate each leaf with a fixed prediction value
- E.g., regression:



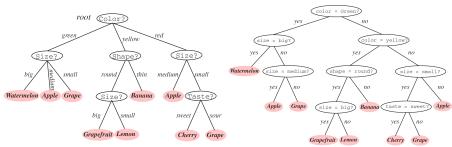




 Key questions: How do we build (learn) the tree? What's the bias/variance tradeoff?

Split factor

- We will focus on binary trees
- This is sufficient: can always convert any tree to binary



from Duda, Hart, and Stork

Regression trees

 \bullet Model corresponding to a tree with M leaves; leaf m corresponds to region R_m which predicts value f_m

$$f(\mathbf{x}) = \sum_{m=1}^{M} f_m \, \mathbb{I}[\mathbf{x} \in R_m]$$

where $\mathbb{I}[A] = 1$ if the predicate A is true and 0 otherwise

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• Given a tree (partition), how do we minimize the squared loss?

$$\min \sum_{i=1}^{n} (f(\mathbf{x}_i) - y_i)^2 \Rightarrow f_m = \frac{1}{|\mathcal{I}_m|} \sum_{i \in \mathcal{I}_m} y_i$$

where $\mathcal{I}_m = \{i: \mathbf{x}_i \in R_m\}$; i.e., f_m is the average value of the training points in R_m .

Regression tree construction

• Goal: find R_1, \ldots, R_M to minimize

$$\sum_{i=1}^{n} \left(\left(\sum_{m=1}^{M} f_m \mathbb{I}[\mathbf{x}_i \in R_m] \right) - y_i \right)^2$$

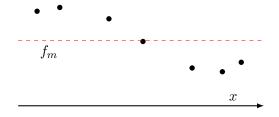
This is not computationally tractable

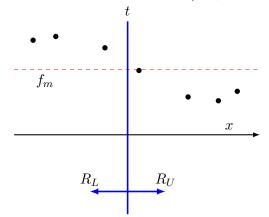
ullet Greedy algorithm instead: consider a split at s along j-th feature,

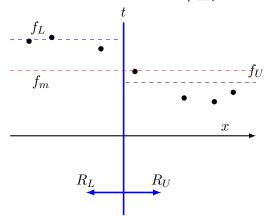
$$R_L(j,s) = \{\mathbf{x} : \phi_j(\mathbf{x}) \le s\}, \quad R_U(j,s) = \{\mathbf{x} : \phi_j(\mathbf{x}) > s\}$$

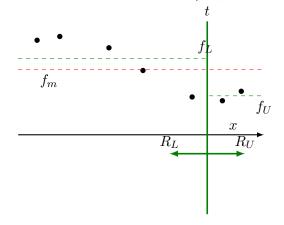
• Cost of the split, assigning $R_L \to f_L$, $R_U \to f_U$:

$$\min_{f_L} \sum_{i: \mathbf{x}_i \in R_L} (y_i - f_L)^2 + \min_{f_U} \sum_{i: \mathbf{x}_i \in R_U} (y_i - f_U)^2$$









Regression tree construction

• So, we need to solve

$$\min_{j,s} \left\{ \min_{f_L} \sum_{i: \mathbf{x}_i \in R_L} (y_i - f_L)^2 + \min_{f_U} \sum_{i: \mathbf{x}_i \in R_U} (y_i - f_U)^2 \right\}$$

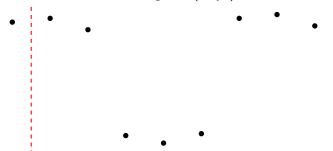
- We already know: for any j, s the f_L , f_U are the averages of the labels in induced R_L , R_U .
- We can exhaustively evaluate all distinct j, s pairs. (What's the running time?)
- ullet Proceed recursively, partitioning R_U and R_L by the same procedure

- Can easily overfit, getting zero training error!
- Idea for limiting model complexity: define notion of "gain", e.g., reduction in loss.
- Now, don't split if the gain is small.
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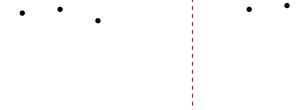
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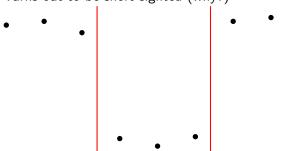


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- Alternative: grow a large tree T_0 , then *prune* to $T \subset T_0$
- Some more notation: |T| is the number of leaves in T, $N_m = |\mathcal{I}_m| = |\{i: \mathbf{x}_i \in R_m\}|$ is the size of a leaf, $f_m = \frac{1}{N_m} \sum_{i \in \mathcal{I}_m} y_i$ is the value in the leaf, and $Q_m(T) = \frac{1}{N_m} \sum_{i \in \mathcal{I}_m} (y_i f_m)^2$ is the leaf error.
- Cost-complexity criterion of tree $T \subset T_0$:

$$C_{\lambda}(T) = \lambda |T| + \sum_{m=1}^{|T|} N_m Q_m(T)$$

Regression tree pruning

$$C_{\lambda}(T) = \lambda |T| + \sum_{m=1}^{|T|} N_m Q_m(T)$$

- T is obtained from T₀ by collapsing some internal nodes (merging multiple leaves)
- For a given $\lambda \geq 0$, there exists a unique $T_{\lambda} = \operatorname{argmin}_{T} C_{\lambda}(T)$
- Weakest link pruning: keep collapsing the internal nodes that produce the *smallest increase* in $\sum_m N_m Q_m(T)$, going from T_0 to a single node.
- Can show: the resulting sequence must contain T_{λ} !
- How do we set λ ?

Classification trees

- What class label \hat{y}_m should we assign to a leaf R_m ?
- Compute fraction of examples from class c in R_m :

$$\hat{p}_{m,c} = \frac{1}{N_m} \sum_{i \in \mathcal{I}_m} \mathbb{I}[y_i = c]$$

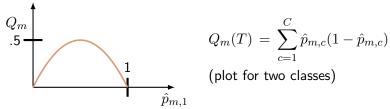
• Want to minimize 0/1 loss, per leaf:

$$\underset{\hat{y}_m}{\operatorname{argmin}} \sum_{i \in \mathcal{I}_m} \mathbb{I}[y_i \neq \hat{y}_m]$$

• Solution: $\hat{y}_m = \operatorname{argmax}_c \hat{p}_{m,c}$

Classification trees: leaf impurity

- In regression trees, squared error is a measure of "leaf impurity"
- In classification, we have a few choices.
- Gini index ("Gini impurity") of leaf m in tree T:



- An alternative Q_m : misclassification rate
- Common practice: use Gini to grow the tree, misclassification rate to prune.

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- Important limitations: hard splits (non-smooth regression), limited to axis-aligned splits, often high variance despite regularization.

Ensembles of models

- So far, we have considered a single model approach: train a model, apply it on test data
- What if we have multiple (different) models?
- **Ensemble** of models: given M models $\{f_1(\mathbf{x}), \dots, f_M(\mathbf{x})\}$, at test time combine them to make a single prediction
- Simplest way to combine: average

$$\widehat{f}(\mathbf{x}) = \frac{1}{M} \sum_{j} f_j(\mathbf{x})$$

When does it help?
 How do we come up with the ensemble?
 Can we do better than just average?

Combining trees

- Deep decision trees have low bias, high variance
- CART pruning may lead to poor bias/variance tradeoff
- Idea: let trees be deep (low bias), average many trees (low variance)

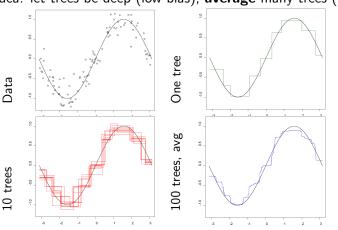
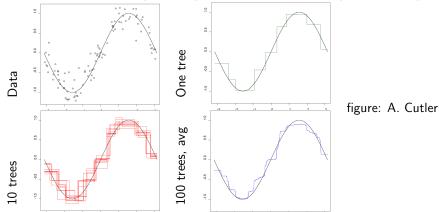


figure: A. Cutler

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• We will now develop a bagging approach (bootstrap aggregation)

Random forests

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- We will obtain diversity by injecting randomness into tree construction

Random forests

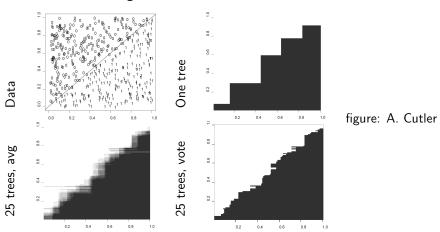
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- Two sources of randomness
- **Bootstrap** sampling: out of n training examples, sample n with replacement some points will appear more than once, some (approx. 37%) will not appear at all

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- Two sources of randomness
- Bootstrap sampling: out of n training examples, sample n with replacement some points will appear more than once, some (approx. 37%) will not appear at all
- Sampling features in each node, when considering splits, only look at a random m < d features.
- Each tree is less likely to overfit
- The "overfitting quirks" of different trees are likely to cancel out in averaging

Classification with random forests

- Tree classifier: compute scores $f_c(\mathbf{x})$, then $h(\mathbf{x}) = \operatorname{argmax}_c f_c(\mathbf{x})$
- Can either average scores, or let trees vote



Random forests summary

- n data points, each with d features
- Build T trees independently (in parallel)
- For each tree: sample n points with replacement (or n' < n without) Grow CART tree; in each node, only look at a random subset of m < d features
 - Do not prune the trees
- To make a prediction: average (regression) or vote (classification)
- Parameters to tune:

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- To make a prediction: average (regression) or vote (classification)
- Parameters to tune: number of trees T; feature set size m; tree depth/min number of points in leaves
- Recommended heuristic values: $m = \sqrt{d}$ for classification, d/3 for regression;
 - min leaf size = 5

Bagging in general

- Instead of trees, could apply bagging to any predictor family
- Power of bagging: variance reduction through averaging
- Typically, benefit is highest with unstable, highly nonlinear predictors (e.g., trees)
- Linear predictors: no benefit from bagging
- Useful property of bagging: "out of bag" (OOB) data in each tree, treat the $\approx \! \! 37\%$ of the examples that didn't make it to the sample as a kind of validation set
- While assembling trees, keep track of OOB accuracy, stop upon seeing plateau

Combining classifiers

- Classifying a point using a decision tree can be seen as a sequence of classifiers, refined as we follow the path to a leaf
- A more general formulation: combine classifiers $h_1(\mathbf{x}), \dots, h_m(\mathbf{x})$

$$H(\mathbf{x}) = \alpha_1 h_1(\mathbf{x}) + \ldots + \alpha_m h_m(\mathbf{x})$$

- α_j is the weight of the vote assigned to classifier h_j
 - Votes should have higher weight for more reliable classifiers
- Prediction (for binary classification):

$$\hat{y}(\mathbf{x}) = \operatorname{sign}(H(\mathbf{x}))$$

• Classifiers h_j can be simple (e.g., based on a single feature)

Greedy assembly of classifier combination

- Consider a family of classifiers \mathcal{H} parametrized by θ .
- Setting θ_1 : minimize the training error

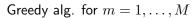
$$\sum_{i=1}^{n} L(h(\mathbf{x}_i; \theta_1), y_i)$$

where L is some surrogate for the 0/1 loss.

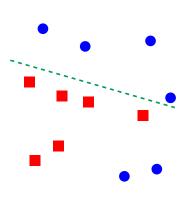
- How do we set θ_2 ?
- We would like to minimize the (surrogate) loss of the combination,

$$\sum_{i=1}^{n} L(H(\mathbf{x}_i), y_i)$$

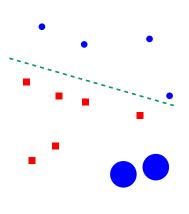
where
$$H(\mathbf{x}) = \operatorname{sign} (\alpha_1 h(\mathbf{x}; \theta_1) + \alpha_2 h(\mathbf{x}; \theta_2))$$



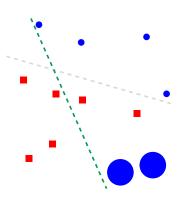
- \bullet Maintain weights $W_i^{(m)}\text{, initially all }1/n$
- ullet Pick a weak classifier h_m minimizing error ϵ_m weighted by $W^{(m-1)}$
- Set $\alpha_m = \frac{1}{2} \log \frac{1 \epsilon_m}{\epsilon_m}$
- Update weights $W_i^{(m)}$ based on mistakes of h_m and on α_m
- Final (strong) classifier $\operatorname{sign}\left(\sum_{m} \alpha_{m} h_{m}(\cdot)\right)$



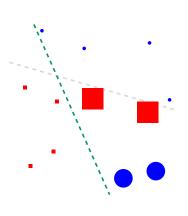
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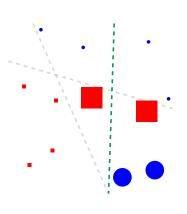
- Maintain weights $W_i^{(m)}$, initially all 1/n
- Pick a weak classifier h_m minimizing error ϵ_m weighted by $W^{(m-1)}$
- Set $\alpha_m = \frac{1}{2} \log \frac{1 \epsilon_m}{\epsilon_m}$
- Update weights $W_i^{(m)}$ based on mistakes of h_m and on α_m
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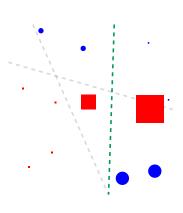
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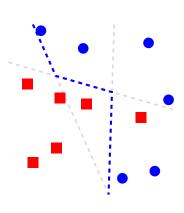
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