Advanced Data Analysis

DATA 71200

Class 9: Naive Bayes and Decision Trees

Naive Bayes

 Uses Bayes rule to predict the probability of an example belonging to a given class

$$P(A \mid B) = \frac{P(B \mid A) \cdot P(A)}{P(B)}$$

A,B = events

P(A|B) = probability of A given B is true

P(B|A) = probability of B given A is true

 $P(A), P(B) \, \mbox{=} \,$ the independent probabilities of A and B

- "Naive" because it examines each feature individually
- Calculate per-class statistics for each feature
 - Bernoulli (binary data) number of times a feature is non-zero for each class
 - Multinomial (count data) average of how many times a feature occurs for each class
 - Gaussian (continuous data) mean and standard deviation of the value of the feature for each class

Naive Bayes

Assumptions

- Independence between features
- Gaussian version assumes normally distributed data with the same variance in each class

Best Practices

- Bernoulli and Multinomial best used on sparse data
- Gaussian version best used on high-dimensional data

Naive Bayes

Parameters

- Alpha (Bernoulli and Multinomial) model complexity
 - The larger the alpha, the more virtual data points that are added for smoothing

Strengths

- Faster to train and predict than linear models
- Understandable training procedure
- Works well on large datasets
 - More efficient on them than linear models

- Coefficients not easily interpreted
- Does not generalize as well as linear models

Learned hierarchy of if/else questions

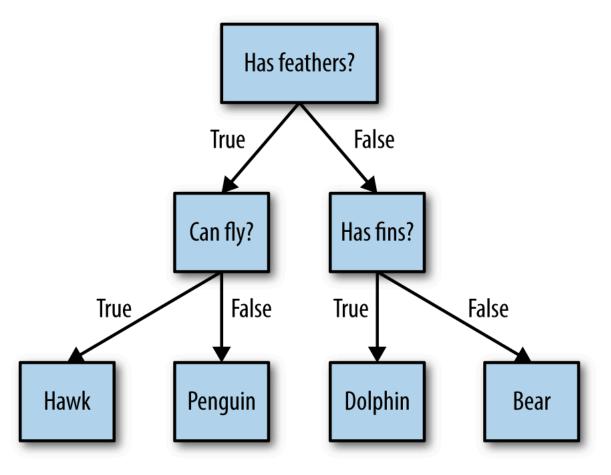


Figure 2-22. A decision tree to distinguish among several animals

Sometimes referred to as CART (Classification and Regression Trees)

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 Algorithm searches through all possible tests to separate the data into classes

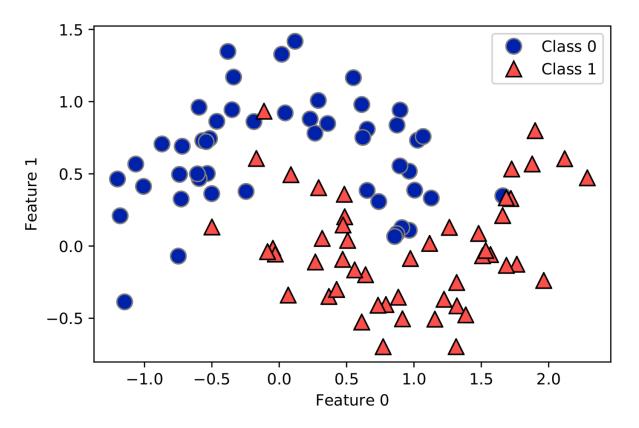
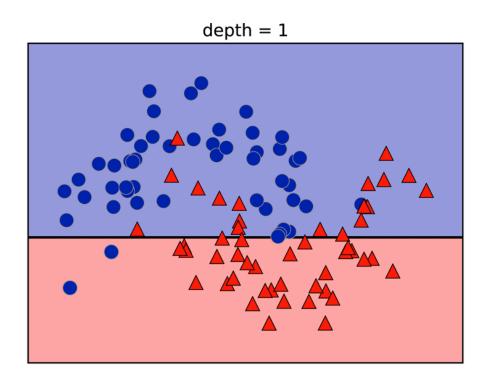


Figure 2-23. Two-moons dataset on which the decision tree will be built

 At each step the algorithm selects the test that provides the best separation of the classes



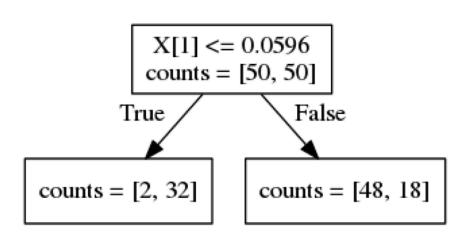
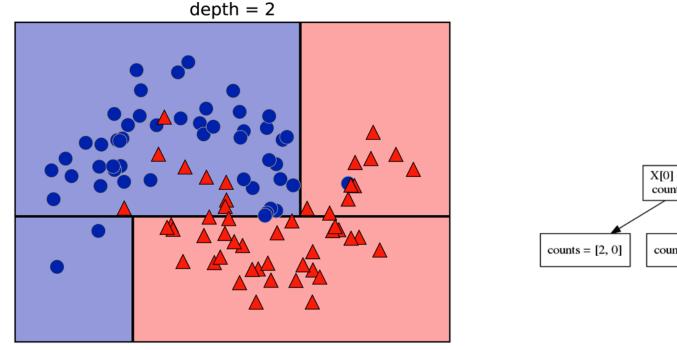


Figure 2-24. Decision boundary of tree with depth 1 (left) and corresponding tree (right)

 At each step the algorithm selects the test that provides the best separation of the classes



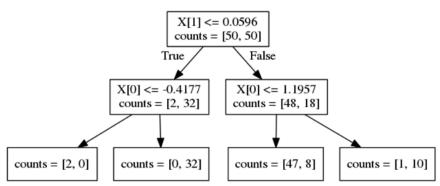
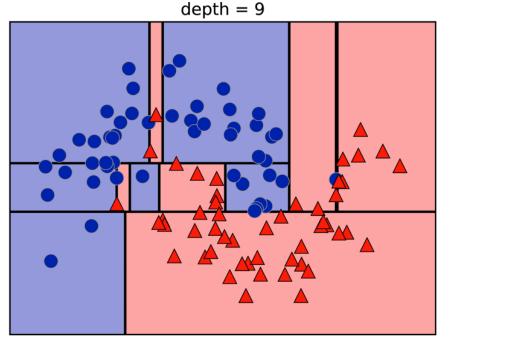


Figure 2-25. Decision boundary of tree with depth 2 (left) and corresponding decision tree (right)

 Partitioning is repeated until each "leaf" contains only a single target (class or regression value) - referred to as pure



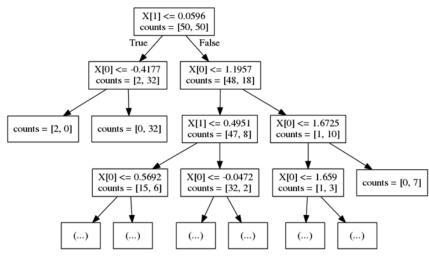
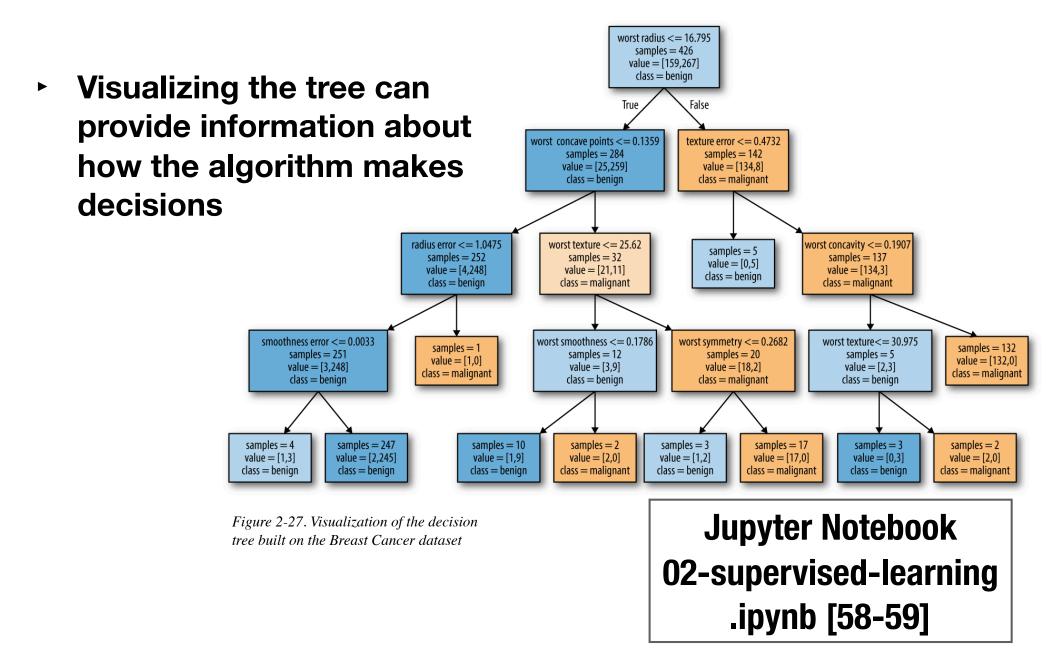


Figure 2-26. Decision boundary of tree with depth 9 (left) and part of the corresponding tree (right); the full tree is quite large and hard to visualize

- Comprehensive tree building leads to overfitting the training data, this can be minimized by
 - Pre-pruning stopping the tree building early
 - Can be achieved by limiting the depth of the tree, number of leaves, or only splitting those with a certain number of points
 - Post-pruning/pruning removing nodes that don't contain much information

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- Feature importance is a summary of how important each feature is in the tree's decision making
- Feature importance isn't class specific

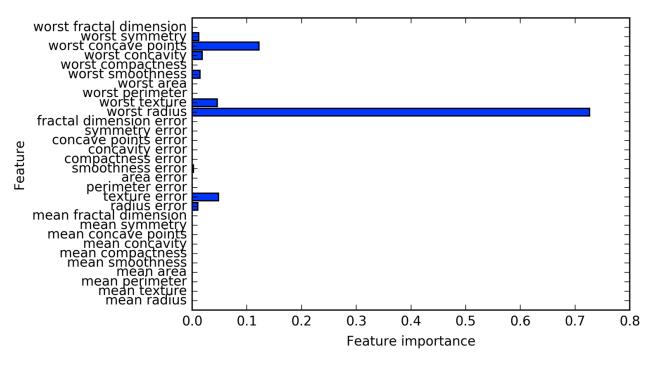
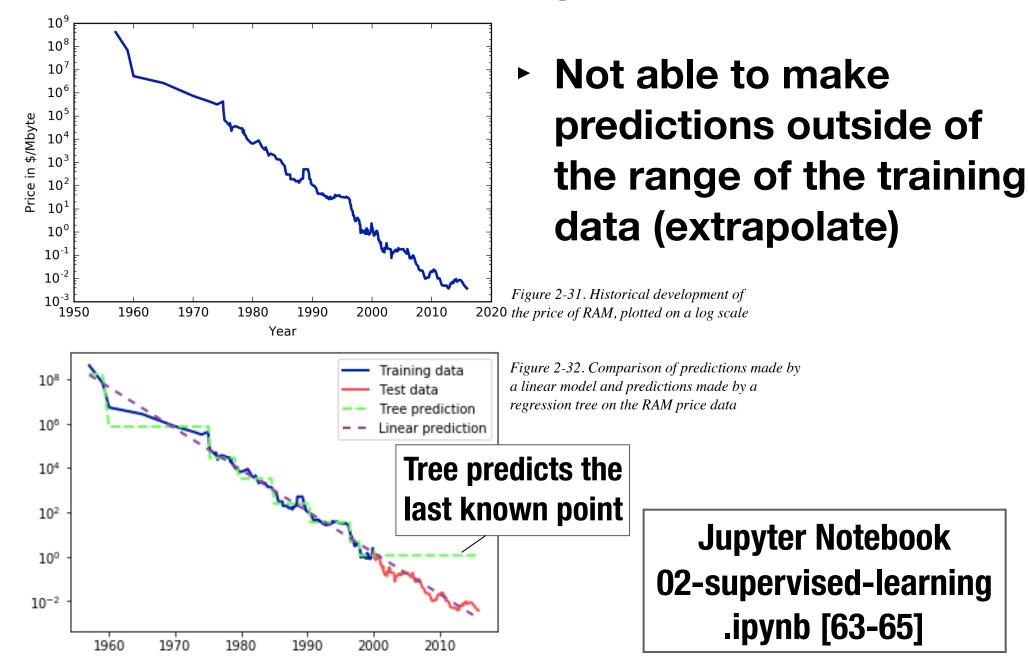


Figure 2-28. Feature importances computed from a decision tree learned on the Breast Cancer dataset

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Decision Trees Regression



Parameters

Maximum depth - for pre-pruning

Assumptions

No assumptions about the distribution of the data

Best Practices

Use of pre-pruning prevents overfitting

Strengths

- Can be visualized, which aids in interpretation
- Invariant to scaling data
- Works well with mixed data (e.g., binary and continuous features)

- Tend to overfit (even with pre-prunning) and thus don't always generalize
- Doesn't work well on high-dimensional sparse data

Random Forests

- Ensemble of slightly different decision trees
 - Averaging the predictions of the trees minimizing overfitting
 - Difference between trees is achieved by randomizing which data points or features are used

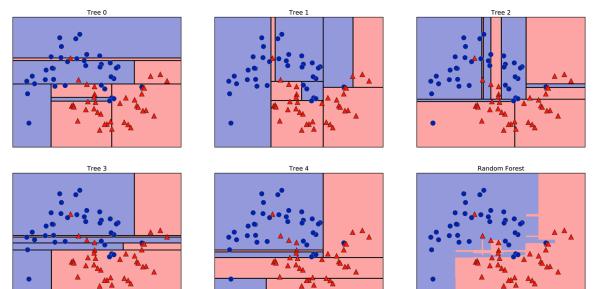


Figure 2-33. Decision boundaries found by five randomized decision trees and the decision boundary obtained by averaging their predicted probabilities

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Random Forests

Feature importance is aggregated across the ensemble of decision trees

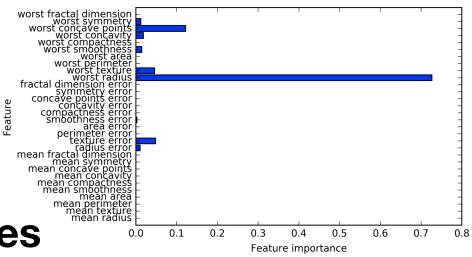


Figure 2-28. Feature importances computed from a decision tree learned on the Breast Cancer dataset

 Typically more informative for random forests than for individual trees

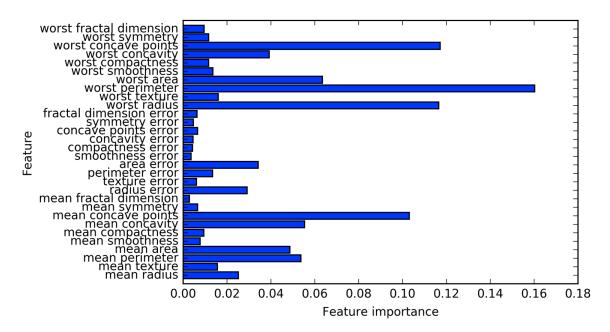


Figure 2-34. Feature importances computed from a random forest that was fit to the Breast Cancer dataset

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Random Forests

Parameters

- Number of estimators larger is always better
- Number of data points number of samples drawn from training dataset
- Maximum features amount of randomness (smaller reduces overfitting)
- Maximum depth for pre-prunning

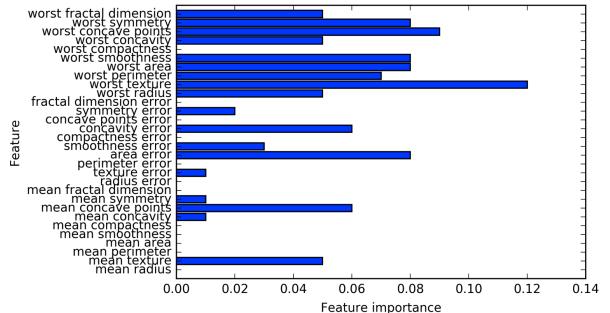
Strength

- Invariant to scaling data
- Works well with mixed data (e.g., binary and continuous features)
- Generalizes better than decision trees

- Harder to interpret than decision trees
- Can be time consuming to train
- Random process can make reproducibility difficult

Gradient Boosted Random Forests

- Series of decision trees where each new tree tries to correct the mistakes of the previous trees using pre-pruning (rather than randomness)
- Trees are typically shallow (weak learners)
- Feature importances tend to be sparser than



random forests

Figure 2-35. Feature importances computed from a gradient boosting classifier that was fit to the Breast Cancer dataset

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Gradient Boosted Random Forests

Best Practices

- Random forests are a good place to start
 - Gradient-boosting can improve accuracy

Parameters

- Number of estimators larger is always better
- Maximum depth for pre-prunning
- Learning rate how much a tree tries to correct the previous mistakes

Strengths

- Typically perform very well
- Invariant to scaling data
- Works well with mixed data (e.g., binary and continuous features)

- Can take a long time to train
- Doesn't work well on high-dimensional sparse data