Classification Methods

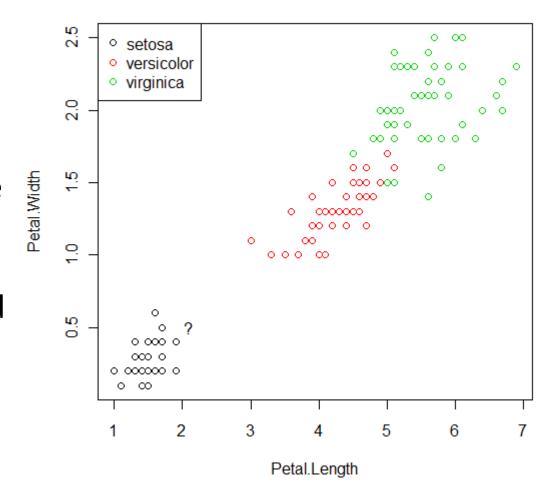
Garrett Dancik, PhD

Classification Methods

- Objective: Identify the class of an individual (e.g., male or female) based on observed features (e.g., gene expression levels)
- Classes: $c_1, c_2, ..., c_m$ Features: $x_1, ..., x_k$
- General Procedure
 - Train the classifier: Using a *training* data set, determine the mapping function $f(x) \rightarrow c$
 - Validation: assess the accuracy of the classifier by applying it to a test data set with known classes
 - Independent validation
 - Leave one out cross validation
 - K-fold cross validation
 - Classification / prediction of target data set

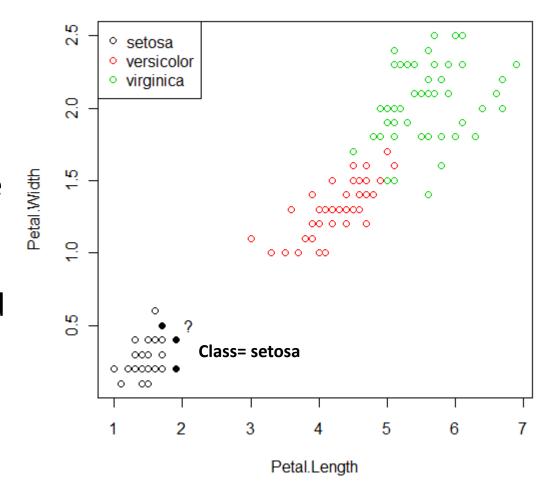
Classification Methods: K-Nearest Neighbors (KNN)

- For a test
 observation A,
 find the distance
 between A and
 every other
 observation in the
 feature space
- Classify the test observation based on the votes of its K nearest neighbors



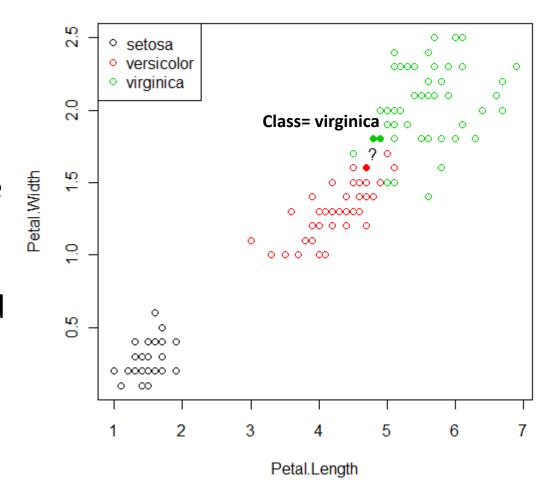
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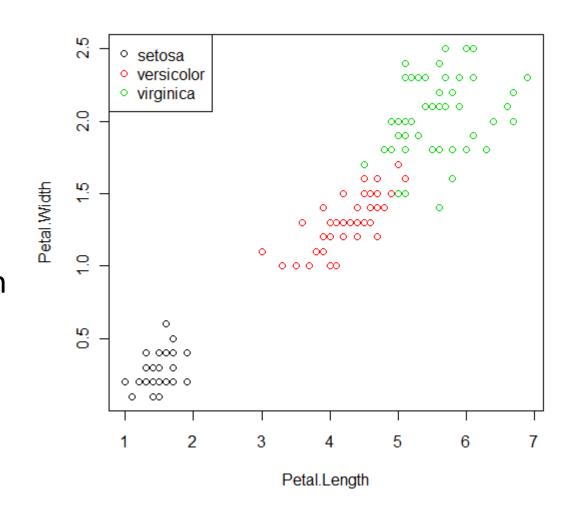


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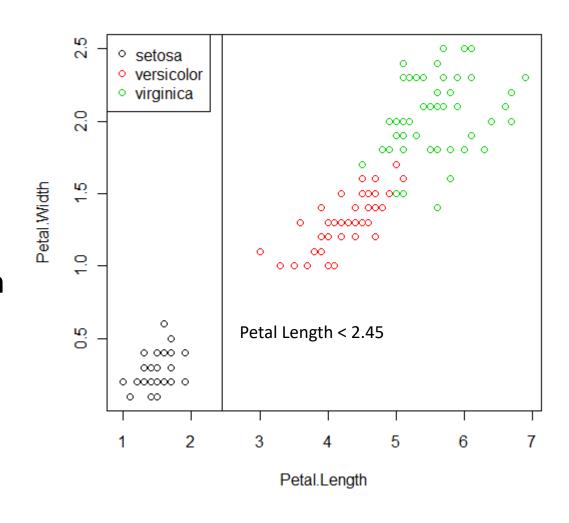
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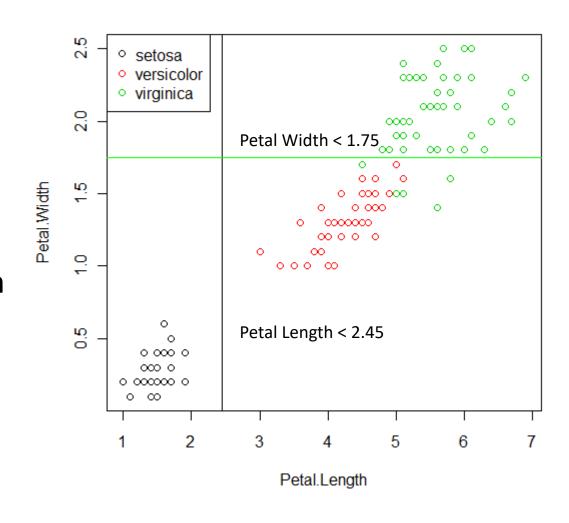
- Create a node by splitting the data according to a feature that optimally splits the data
- Repeat on data subsets until a stopping criterion is met
- Each leaf corresponds to a class

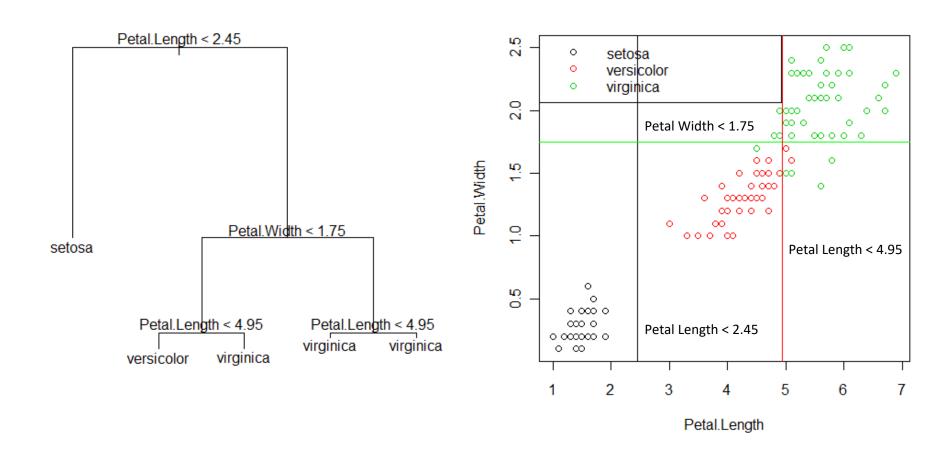


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Note: DT are known to overfit data. However more rubust methods such as Random Forests can be used

Classification Methods: Naïve Bayes (NB)

Based on Bayes' theorem that relates conditional probabilities

$$p(C|x_1,...,x_n) \propto p(x_1,...,x_n|C)p(C)$$

 Naïve Bayes assumes independence of features, so that

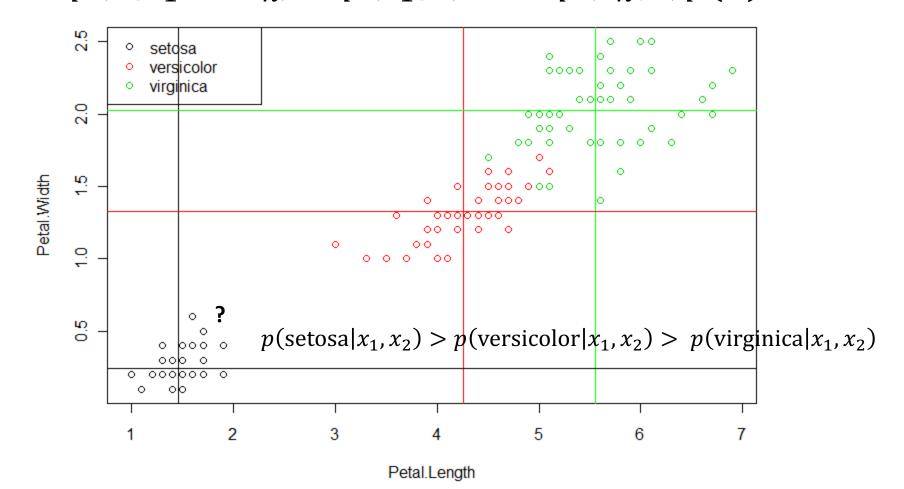
$$p(x_1, ..., x_n | C) \propto p(x_1 | C) \times ... \times p(x_n | C) p(C)$$

• For quantitative features, calculate by treating $p(x|\mathcal{C}) \sim N(\mu_{x},\sigma_{x})$

• Select the class C that maximizes $p(C|x_1,...,x_n) \propto p(x_1|C) \times \cdots \times p(x_n|C)p(C)$

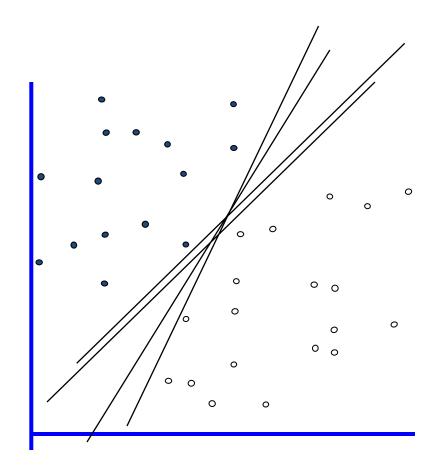
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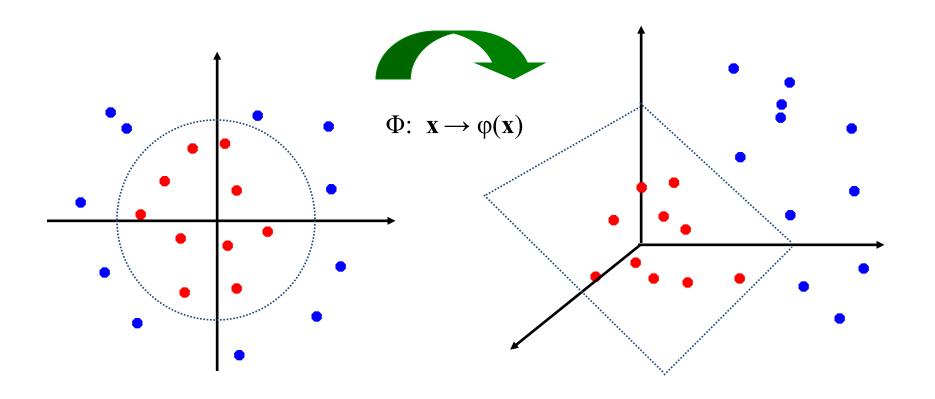


Classification Methods: Support Vector Machines (SVM)

- Find the optimum hyperplane that linearly separates the classes
- If classes are not linearly separable, map the data into a higher dimensional space through the use of a kernel function



Classification Methods: Support Vector Machines (SVM)



Caveats and strategies

Validation

- Overfitting is often a problem: a classifier can perform very well on a training data set but may not generalize to additional data sets
- Validation on independent data sets are ideal
- Cross-validation is useful when data is limited

Basic Strategy

- Use cross-validation to select
 - The number of features (e.g., probes/genes)
 - Optimal parameters for classification model (e.g, value of k in knn)