Instance-Based Learning

www.biostat.wisc.edu/~dpage/cs760/

Goals for the lecture

you should understand the following concepts

- *k*-NN classification
- k-NN regression
- edited nearest neighbor
- k-d trees for nearest neighbor identification

Nearest-neighbor classification

learning task

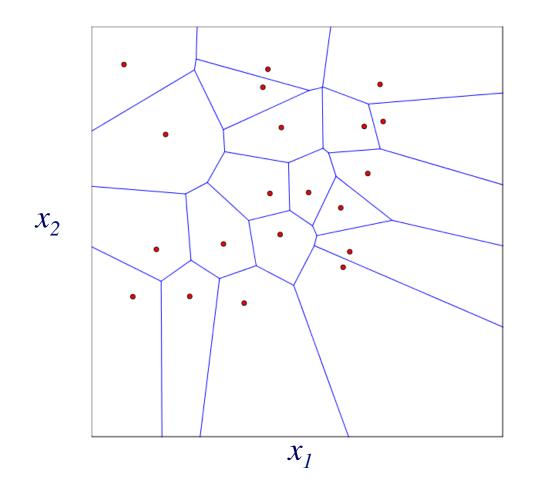
given a training set (x₁,y₁)...(x_n,y_n), do nothing (it's sometimes called a *lazy learner*)

classification task

- **given**: an instance x_q to classify
- find the training-set instance x_i that is most similar to x_q
- return the class value y_i

The decision regions for nearestneighbor classification

Voronoi Diagram: Each polyhedron indicates the region of feature space that is in the nearest neighborhood of each training instance



k-nearest-neighbor classification

classification task

- **given**: an instance x_q to classify
- find the k training-set instances (x₁,y₁)...(x_k,y_k) that are most similar to x_q
- return the class value

$$\hat{y} \leftarrow \underset{v \in \text{values}(Y)}{\operatorname{arg\,max}} \sum_{i=1}^{k} \delta(v, y_i) \qquad \qquad \delta(a, b) = \begin{cases} 1 \text{ if } a = b \\ 0 \text{ otherwise} \end{cases}$$

(i.e. return the class that the plurality of the neighbors have)

How can we determine similarity/distance

suppose all features are nominal (discrete)

 Hamming distance: count the number of features for which two instances differ

suppose all features are continuous

• Euclidean distance:

$$d(\boldsymbol{x}_i, \boldsymbol{x}_j) = \sqrt{\sum_{f} (x_{if} - x_{jf})^2} \quad \text{where } x_{if} \text{ represents the } f^{th} \text{ feature of } \boldsymbol{x}_i$$

• Could also use Manhattan distance: sum the differences in feature values – continuous analog to Hamming distance

How can we determine similarity/distance

if we have a mix of discrete/continuous features:

$$d(\boldsymbol{x}_{i}, \boldsymbol{x}_{j}) = \sum_{f} \begin{cases} \left| x_{if} - x_{jf} \right| & \text{if } f \text{ is continuous} \\ 1 - \delta(x_{if}, x_{jf}) & \text{if } f \text{ is discrete} \end{cases}$$

 If all feature are of equal importance, want to apply to continuous features some type of normalization (values range 0 to 1) or standardization (values distributed according to standard normal)

k-nearest-neighbor regression

learning task

• given a training set $(x_1, y_1)...(x_n, y_n)$, do nothing

prediction task

- **given**: an instance x_q to make a prediction for
- find the *k* training-set instances (*x*₁, *y*₁)...(*x*_k, *y*_k) that are most similar to *x*_q
- return the value

$$\hat{y} \leftarrow \frac{1}{k} \sum_{i=1}^{k} y_i$$

Distance-weighted nearest neighbor

We can have instances contribute to a prediction according to their distance from x_a

classification:

$$\hat{y} \leftarrow \underset{v \in \text{values}(Y)}{\operatorname{arg\,max}} \sum_{i=1}^{k} w_i \ \delta(v, y_i)$$

 $w_i = \frac{1}{d(x_a, x_i)^2}$

regression:

$$\hat{y} \leftarrow \frac{\sum_{i=1}^{k} w_i y_i}{\sum_{i=1}^{k} w_i}$$

Speeding up k-NN

- *k*-NN is a "lazy" learning algorithm does virtually nothing at training time
- but classification/prediction time can be costly when the training set is large
- two general strategies for alleviating this weakness
 - don't retain every training instance (edited nearest neighbor)
 - use a smart data structure to look up nearest neighbors (e.g. a k-d tree)

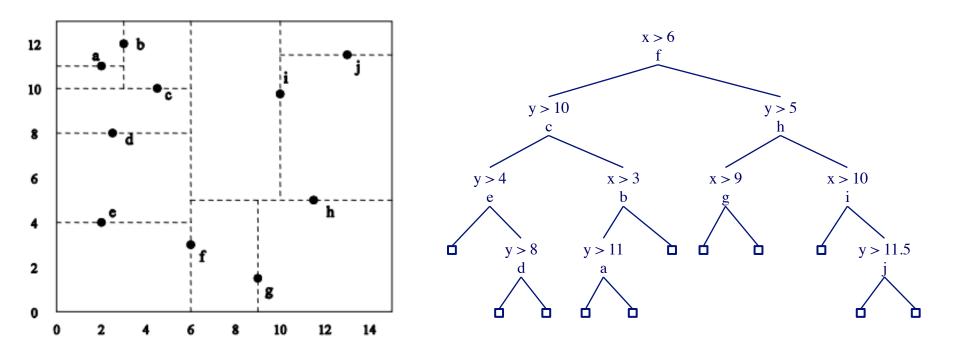
Edited instance-based learning

- select a subset of the instances that still provide accurate classifications
- *incremental deletion*
 start with all training instances in memory
 for each training instance (*x_i*, *y_i*)
 if other training instances provide correct classification for (*x_i*, *y_i*)
 delete it from the memory
- incremental growth
 start with an empty memory
 for each training instance (x_i, y_i)
 if other training instances in memory don't correctly classify (x_i, y_i)
 - add it to the memory

k-d trees

a *k-d tree* is similar to a decision tree except that each internal node

- stores one instance
- splits on the median value of the feature having the highest variance



Finding nearest neighbors with a k-d tree

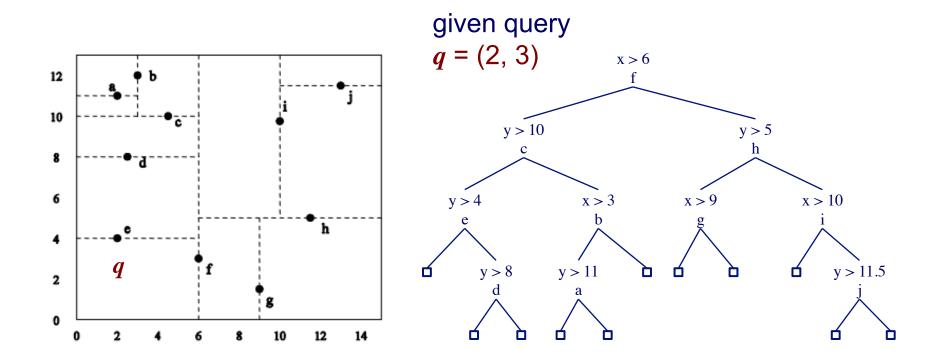
- use branch-and-bound search
- priority queue stores
 - nodes considered
 - lower bound on their distance to query instance
- lower bound given by distance using a single feature
- average case: $O(\log_2 n)$
- worst case: O(n)

Finding nearest neighbors in a k-d tree

NearestNeighbor(instance q)

 $PQ = \{\}$ // minimizing priority queue best_dist = ∞ // smallest distance seen so far PQ.push(root, 0) while PQ is not empty (node, bound) = PQ.pop();if (bound \geq best dist) // bound is best bound on all PQ return best_node.instance // so nearest neighbor found dist = distance(q, node. instance) if (dist < best_dist) best dist = dist best node = node if (q[node.feature] - node.threshold > 0)PQ.push(node.left, *q*[node.feature] – node.threshold) PQ.push(node.right, 0) else PQ.push(node.left, 0) PQ.push(node.right, node. threshold - q [node.feature]) return best node. instance

k-d tree example (Manhattan Distance)



distance	best distance	best node	priority queue
	∞		(f, 0)
4.0	4.0	f	(c, 0) (h, 4)
10.0	4.0	f	(e, 0) (h, 4) (b, 7)
1.0	1.0	е	(d, 1) (h, 4) (b, 7)

Strengths of instance-based learning

- simple to implement
- "training" is very efficient
- adapts well to on-line learning
- robust to noisy training data (when k > 1)
- often works well in practice

Limitations of instance-based learning

- sensitive to range of feature values
- sensitive to irrelevant and correlated features, although...
 - there are variants that learn weights for different features
 - later we'll talk about *feature selection* methods
- classification can be inefficient, although edited methods and k-d trees can help alleviate this weakness
- doesn't provide much insight into problem domain because there is no explicit model